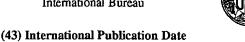
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(54) Title: METHODS FOR PROPAGATING ADENOVIRUS AND VIRUS PRODUCED THEREBY

(57) Abstract: Various methods for propagating and rescuing multiple serotypes of replication-defective adenovirus in a single adenoviral E1-complementing cell line are disclosed. Typically, replication-defective adenovirus vectors propagate only in cell lines which express E1 proteins of the same serotype or subgroup as the vector. The disclosed methods offer the ability to propagate vectors derived from multiple adenoviral serotypes in a single production cell line which expresses E1 proteins from a single serotype. Propagation in this manner is accomplished by providing all or a portion of an E4 region in cis within the genome of the replicationdefective adenovirus. The added E4 region or portion thereof is cloned from a virus of the same or highly similar serotype as that of the E1 gene product(s) of the complementing cell line. Interaction between the expressed E1 of the cell line and the heterologous E4 of the replication-defective adenoviral vectors enables their propagation and rescue. The invention bypasses a need in the art to customize specific cell lines to the serotype or subgroup of the adenoviral vector being propagated and enables one to easily and rapidly develop alternative adenoviral serotypes as gene delivery vectors for use as vaccines or as a critical component in gene therapy.



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Electronic data base consulted during the international search (name of data base and, where practicable, search terms used) PubMed									
C. DOC	UMENTS CONSIDERED TO BE RELEVANT								
Category *	Citation of document, with indication, where a	ppropriate, of the relevant passages	Relevant to claim No.						
х - Y	US 5,837,511 A (FALCK-PEDERSEN et al.) 17 No lines 47-55, 62-67, column 8, lines 6-18, column 9, 35-67, column 16, lines 23-49		1, 2, 4-10, 12, 13, 17- 21, 24, 27-37, 43-50, 56-65, 71-78						
			3, 11, 14-16, 22, 23, 25, 26, 38-42, 51-55, 66-70, 79-83						
Y	US 6,200,798 B1 (YEH et al.) 13 March 2001 (13.0	3, 14-16, 22, 23, 25, 26, 38-42, 51-55, 66- 70, 79-83							
. <b>Y</b>	US 6,391,612 B1 (BRUDER et al.) 21 May 2002 (2 column 6, lines 1-10	1.05.2002), column 5, lines 60-67,	11						
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Further	documents are listed in the continuation of Box C.	See patent family annex.							
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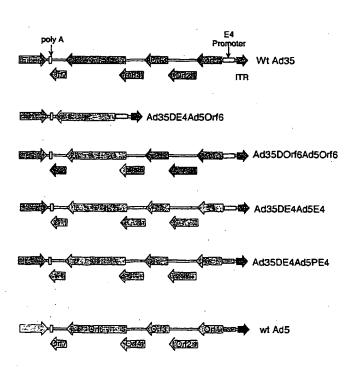
- (71) Applicant (for all designated States except US): MERCK & CO., INC. [US/US]; 126 East Lincoln Avenue, Rahway, NJ 07065-0907 (US).
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[Continued on next page]

#### (54) Title: METHODS FOR PROPAGATING ADENOVIRUS AND VIRUS PRODUCED THEREBY



Various methods for propagating and rescuing multiple serotypes of replication-defective adenovirus in a single adenoviral E1-complementing cell line are disclosed. Typically, replication-defective adenovirus vectors propagate only in cell lines which express E1 proteins of the same serotype or subgroup as the vector. The disclosed methods offer the ability to propagate vectors derived from multiple adenoviral serotypes in a single production cell line which expresses E1 proteins from a single serotype. Propagation in this manner is accomplished by providing all or a portion of an E4 region in cis within the genome of the replication-defective adenovirus. The added E4 region or portion thereof is cloned from a virus of the same or highly similar serotype as that of the E1 gene product(s) of the complementing cell line. Interaction between the expressed E1 of the cell line and the heterologous E4 of the replication-defective adenoviral vectors enables their propagation and rescue. The invention bypasses a need in the art to customize specific cell lines to the serotype or subgroup of the adenoviral vector being propagated and enables one to easily and rapidly develop alternative adenoviral serotypes as gene delivery vectors for use as vaccines or as a critical component in gene therapy.

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## TTILE OF THE INVENTION

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METHODS FOR PROPAGATING ADENOVIRUS AND VIRUS PRODUCED THEREBY

## CROSS-REFERENCE TO RELATED APPLICATIONS

The present application claims the benefit of application serial nos. 60/458,825, filed March 28, 2003; 60/455,312, filed March 17, 2003; 60/455,234, filed March 17, 2003; and 60/405,182, filed August 22, 2002.

#### FIELD OF THE INVENTION

The present invention concerns various methods to propagate and rescue multiple serotypes of replication-defective adenovirus in a single adenoviral E1-complementing cell line. Typically, replication-defective adenovirus vectors propagate only in cell lines which express E1 proteins of the same serotype or subgroup as the vector. The methods disclosed herein offer the ability to propagate vectors derived from multiple serotypes in a single cell line expressing E1 proteins from a single serotype. Such propagation of a wide range of vectors in one cell line is accomplished by providing all or a portion of an E4 region in cis within the genome of the replication-defective adenovirus. The added E4 region or portion thereof is cloned from a virus of the same or highly similar serotype as that of the E1 gene product(s) of the complementing cell line. Interaction between the E1 gene products of the cell line and the heterologous E4 gene products of the replication-defective adenoviral vector enables the propagation and rescue of the recombinant replication-defective adenovirus vectors. The invention, therefore, bypasses an existing need in the art to customize complementing cell lines to the specific serotype or subgroup of the adenoviral vector being propagated or, alternatively, to have to transfect a cell line with an E4 region and then regulate the expression in trans of the E4 region within the E1 complementing cell line.

## BACKGROUND OF THE INVENTION

Beginning with the first human adenoviruses (Ads) isolated over four decades ago (Rowe et al., Proc. Soc. Exp. Biol. Med., 84:570-579, 1953), over 100 distinct serotypes of adenovirus have been isolated which infect various mammalian species, 51 of which are of human origin (Straus, Adenovirus infections in humans. In The Adenoviruses. 451-498, 1984; Hierholzer et al., J. Infect. Dis., 158: 804-813, 1988; Schnurr and Dondero, Intervirology., 36: 79-83, 1993; Jong et al., J Clin Microbiol., 37:3940-3945:1999). The human serotypes have been categorised into six subgenera (A-F) based on a number of biological, chemical, immunological and structural criteria; criteria which include hemagglutination properties of rat

and rhesus monkey erythrocytes, DNA homology, restriction enzyme cleavage patterns, percentage of G+C content and oncogenicity (Straus, Adenovirus infections in humans. In *The Adenoviruses*. 451-498, 1984; Horwitz, Adenoviridae and their replication, *In Virology*: 1679-172, 1990).

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Deletion of an essential E1 region common to the various adenovirus serotypes has enabled the use of adenovirus vectors as gene transfer vectors for vaccine and gene therapy purposes. Resultant replication-defective vectors are propagated in cell lines that provide the deleted E1 gene products in trans. Supplementation of the essential E1 gene products in trans in this manner works well when the E1 gene products are from the same or a highly similar serotype. As such, E1-deleted group C serotypes (Ad1, Ad2, Ad5 and Ad6) grow well in 293 or PER.C6 cells which contain and express the Ad5 E1 region. In contrast, E1-deleted serotypes other than group C, for example those from subgroups A, B, D, E, and F (e.g., Ad3, Ad4, and Ad7 to Ad51), do not replicate efficiently in 293 or PER.C6 cells. The Ad5 E1 sequences in 293 and PER.C6 cells do not fully complement the replication of these alternative serotypes. This presents a challenge due to the fact that the most characterized and studied complementing cell lines available for growth and propagation of adenovirus are based on E1 sequence from adenovirus serotype 5.

This inability to fully complement the replication of serotypes other than group C adenovirus in Ad5 E1 complementing cell lines has been attributed to the inability of Ad5 (group C) E1b 55K gene product to functionally interact with the E4 gene products of non-group C serotypes. While the interaction is conserved within members of the same subgroup, it is not well conserved between subgroups.

Hence, cell lines expressing both Ad5 E1 and ORF6 were generated and proved useful in complementing alternative adenovirus serotypes; see, e.g., Abrahamsen et al., 1997 J. Virol. 8946-8951. Such incorporation of E4 (or ORF6) into Ad 5 complementing cell lines as was done in Abrahamsen et al., supra, is known.

U.S. Patent No. 5,849,561 discloses complementation of an E1-deleted non-group C adenovirus vector in an Ad5-E1 complementing cell line which also expresses portions of the Ad5-E4 gene.

U.S. Patent No. 6,127,175, issued to Vigne, *et al.*, discloses a stably transfected mammalian cell line which expresses a portion of the E4 region of adenovirus, preferably ORF6 or ORF6/7. Such a cell line is useful for complementation of recombinant Ad genomes deficient in the E4 region.

European Application EP 1 054 064 A1 discloses recombinant, replication deficient adenovirus 35 (Ad35) vectors and cell lines which complement in trans the growth of

these vectors. A cell line which expresses Ad5E1A and E2A genes (PER.C6) was shown to complement an Ad35-E1 deleted vector upon co-expression of Ad35-E1B proteins.

U.S. Patent No. 6,270,996, issued to Wilson, et al., discloses E1/E4 deleted adenovirus vectors and E1/E4(ORF6) cell lines which complement in trans virus growth without resulting in cell toxicity.

U.S. Patent No. 6,202,060, issued to Mehtali, et al., discloses adenoviral vectors wherein portions of the early genes are under control of an inducible promoter. The '060 patent also discloses complementing cell lines which may be used in tandem with these Ad vectors.

The generation of serotype-specific cell lines providing a complementing serotype-specific E1 gene product(s) in trans is known as well.

Although Ad5-based vectors have been used extensively in a number of gene therapy trials, there may be limitations on the use of Ad5 and other group C adenoviral vectors due to preexisting immunity in the general population due to natural infection. Ad5 and other group C members tend to be among the most seroprevalent serotypes. Immunity to existing vectors may develop as a result of exposure to the vector during treatment. These types of preexisting or developed immunity to seroprevalent gene delivery vectors may limit the effectiveness of gene therapy or vaccination efforts. Alternative adenovirus serotypes, thus, constitute very important targets in the pursuit of gene delivery systems capable of evading the host immune response.

There remains both a practical and commercial need for an adenovirus-based vaccine and/or gene therapy delivery system which allows for the production of multiple serotype recombinant adenovirus vectors in a single source complementing mammalian cell line. The present invention addresses and overcomes this deficiency in the art by disclosing novel methods for propagating multiple serotype recombinant Ad vectors in a single complementing cell line where the required serotype-specific sequences are provided *in cis*.

## SUMMARY OF THE INVENTION

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The present invention relates to an enhanced means for propagating replication-defective adenovirus in an E1-complementing cell line(s) where the E1 gene product(s) being expressed is not native to the adenovirus being propagated. The method is based on Applicants' finding that supply, in cis, of a nucleic acid sequence encoding all or a portion of a heterologous adenoviral E4 region which is native to a virus of the same or highly similar serotype as the E1 gene product(s) of the complementing cell line enables the growth of adenoviral vectors of varying serotype in any single complementing cell line, despite the fact the cell line is not customized for the particular serotype of vector being propagated. This is of particular

importance given that existing and settled adenoviral E1-complementing cell lines (such as PER.C6™ and 293) are based on one of the most prominent adenovirus serotypes (Ad5) and are not suited for the large-scale propagation and rescue of alternative serotypes.

The basic steps involved in the propagation of adenoviral vectors in accordance with the methods of the instant invention are as follows: First, all or a portion of a heterologous adenoviral E4 region comprising nucleic acid sequence encoding at least open reading frame 6 (ORF6) is inserted into a replication-defective adenoviral vector. By "heterologous", Applicants mean that the nucleic acid sequence is not native to the viral vector being propagated, i.e., not normally present within a virus of the same or highly similar serotype. As will be described, the adenoviral E4 region or portion thereof can be either a nucleic acid sequence encoding ORF 6 or any larger portion of the E4 region, and includes nucleic acid comprising the complete E4 region with E4 promoter. The region into which the nucleic acid is incorporated is not limited, i.e., the insertion can be made into the complete E4 region with E4 promoter or into a smaller portion narrowing into the ORF6 region. Alternatively, the heterologous E4 region or portion thereof can be inserted into different areas of the genome such as the E1 or E3 regions. Further, the native E4 region or portion thereof can be deleted and replaced, or left intact. This is not deemed a critical element of the instant invention. What is a critical element is that the heterologous E4 region or portion thereof being inserted is native to a virus of the same or highly similar serotype as the E1 gene product(s) expressed by the complementing cell line.

Following the modification of the adenoviral vector of interest, the recombinant adenovirus is then introduced into an adenoviral E1-complementing cell line and allowed to propagate. The adenovirus is subsequently harvested and rescued from the complementing cell line.

The resultant virus can be studied and used in various gene therapy and vaccine efforts. The virus, therefore, forms an important aspect of the instant invention.

## BRIEF DESCRIPTION OF THE DRAWINGS

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FIGURE 1 illustrates a transcription map for adenovirus serotype 5. The linear genome is divided into 100 map units as well as into r- and l- strands which designate the direction of transcription. Early transcription units are designated with an E and are active prior to viral DNA replication. Late transcription units are designated with and L and are active primarily after DNA replication. Promoters are represented as brackets and polyadenylation sites as arrowheads. The tripartite leader is designated 1, 2, and 3.

FIGURES 2A-1 through 2A-10 illustrate the nucleic acid sequence for the wild-type adenovirus 35 (SEQ ID NO: 1) utilized in the Examples.

FIGURE 3 illustrates the homologous recombination scheme utilized to recover pAd35 $\Delta$ E1.

FIGURE 4 illustrates the various configurations of the E4 regions (or portions) within the alternative serotype recombinants.

FIGURE 5 illustrates the homologous recombination scheme utilized to recover pAd35 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

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FIGURE 6 illustrates the nucleic acid sequence encoding the gag expression cassette (SEQ ID NO: 2). The various regions of the figure are as follows: (1) a first underlined segment of nucleic acid sequence encoding the immediate early gene promoter region from human cytomegalovirus; (2) a first segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; (3) a region in caps which contains the coding sequence of HIV-1 gag; (4) a second segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; and (5) a second underlined segment, this segment containing nucleic acid sequence encoding a bovine growth hormone polyadenylation signal sequence.

FIGURE 7 illustrates the nucleic acid sequence encoding the SEAP expression cassette (SEQ ID NO: 3). The various regions of the figure are as follows: (1) a first underlined segment of nucleic acid sequence encoding the immediate early gene promoter region from human cytomegalovirus; (2) a first segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; (3) a region in caps which contains the coding sequence of the human placental SEAP gene; (4) a second segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; and (5) a second underlined segment, this segment containing nucleic acid sequence encoding a bovine growth hormone polyadenylation signal sequence.

FIGURE 8 illustrates *in vivo* expression of SEAP in C3H/HeN mice using 10<sup>10</sup> vp doses of Ad35 vectors. This experiment was designed to address any effects of E3 deletion. The vectors were injected intramuscularly and the levels of SEAP expression were determined from the serum samples. Shown are geometric means for each cohort of 5 mice.

FIGURE 9 illustrates in vivo expression of SEAP in C3H/HeN mice using 10^10 vp doses of Ad35 vectors. This experiment was designed to address any effects of Ad5 sequence insertion into the Ad35 genome. The vectors were injected intramuscularly and the levels of SEAP expression were determined from the serum samples. Two extra cohorts received 10^10 vp and 10^9 vp of Ad5 vector. Shown are geometric means for each cohort of 5 mice.

FIGURES 10A-B illustrate *in vivo* SEAP expression using MRKAd5-based (A) and Ad35ΔE1ΔE4Ad5Orf6-based (B) vector in rhesus macaques. Shown are the serum antigen

levels for individual monkeys following a single intramuscular (i.m.) injection of 10<sup>11</sup> vp MRKAd5SEAP (filled circles), 10<sup>9</sup> vp MRKAd5SEAP (open boxes) or 10<sup>11</sup> vp Ad35ΔE1SEAPΔE4Ad5Orf6.

FIGURE 11 illustrates in vivo SEAP expression in African green monkeys using Ad5- and Ad35-based vectors. Shown are the antigen levels for each animal in serum samples collected two days after the treatment.

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FIGURE 12 illustrates the homologous recombination scheme utilized to recover pAd24 $\Delta$ E1.

FIGURE 13 illustrates the homologous recombination scheme utilized to recover pAd24ΔE1Ad5Orf6.

FIGURE 14 illustrates the configuration of E4 regions in the Ad24 recombinants generated.

FIGURE 15 illustrates the growth kinetics of the Ad24-based vectors in PER.C6 cells.

FIGURES 16A-1 through 16A-10 illustrate the nucleic acid sequence for wild-type adenovirus serotype 24 (SEQ ID NO: 5). The ATCC product number for Ad24 is VR-259.

FIGURE 17 illustrates, in tabular format, gag-specific T cell responses in monkeys immunized with MRKAd5-HIVgag and Ad24 HIV vectors. Shown are the numbers of spot-forming cells per million PBMC following incubation in the absence (mock) or presence of Gag peptide pool. The pool consisted of 20-aa peptide overlapping by 10 aa and encompassing the entire gag sequence.

FIGURE 18 illustrates, in tabular format, the characterization of the gag-specific T cells in monkeys immunized with 10^11 vp of MRKAd5-HIV1gag and Ad24ΔE1gagΔOrf6Ad5Orf6. Shown are the percentages of CD3+ T cells that are either gag-specific CD4+ or gag-specific CD8+ cells. These values were corrected for mock values (<0.03%).

FIGURE 19 illustrates individual anti-p24 titers (in mMU/mL) in macaques immunized with gag-expressing adenovirus vectors.

FIGURE 20 illustrates in vivo expression of SEAP in C3H/HeN mice using 10^10 vp doses of Ad24 vectors. The vectors were injected intramuscularly and the levels of SEAP expression were determined from the serum samples. Two extra cohorts received 10^10 vp and 10^9 vp of Ad5 vector. Shown are geometric means for each cohort of 5 mice.

FIGURE 21 illustrates *in vivo* SEAP expression using MRKAd5 and Ad24 vectors in rhesus macaques. Shown are the geometric means of the SEAP levels for cohorts of 3 monkeys. In bars are the standard errors of the geometric means.

FIGURE 22 illustrates a homologous recombination scheme to be utilized to recover pAd24 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

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FIGURE 23 illustrates gag-specific T cell responses in rhesus macaques immunized following a heterologous Ad5/Ad6 prime-Ad24 boost regimen. a: Mock, no peptide: gag, 20-mer peptide pool encompassing entire gag sequence; b: Peak response after 2 or 3 doses of the priming vaccine; c: 3 wks prior to boost; d: 4 wks after boost; e: ND, not determined.

FIGURE 24 illustrates, in tabular format, the percentages of CD3<sup>+</sup> T lymphocytes that are gag-specific CD8<sup>+</sup> cells or gag-specific CD4<sup>+</sup> cells determined after the Ad24 Boost Immunization (wk 60). Numbers reflect the percentages of circulating CD3+ lymphocytes that are either gag-specific CD4+ or gag-specific CD8+ cells. Mock values (equal to or less than 0.01%) have been subtracted.

FIGURE 25 illustrates gag-specific T cell responses in rhesus macaques immunized following a heterologous Ad 24 prime-Ad5 boost regimen. a: Mock, no peptide: gag, 20-mer peptide pool encompassing entire gag sequence; b: Peak response after 2 doses of the priming vaccine; c: Wk 24; d: 4 wks after boost; e: ND, not determined.

FIGURE 26 illustrates the homologous recombination scheme utilized to recover pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

FIGURE 27 illustrates the homologous recombination scheme utilized to recover pMRKAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

FIGURES 28A-1 to 28A-9 illustrate a nucleic acid sequence for wild-type adenovirus serotype 34 (SEQ ID NO: 12). The ATCC product number for Ad34 is VR-716.

FIGURE 29 illustrates the time course of SEAP expression using MRKAd5 and

FIGURE 29 illustrates the time course of SEAP expression using MRKADS and Ad34 vectors in rhesus macaques. Data represent cohort geometric means.

FIGURE 30 illustrates, in tabular format, T cell responses induced using MRKAd5 and Ad34 vectors expressing HIV-1 gag. Data are expressed in numbers of spot-forming cells per million PBMC (SFC/10^6 PBMC). "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

FIGURE 31 illustrates, in tabular format, the levels of CD4+ and CD8+ Gag-specific T cells in Ad34-immunized macaques at week 12. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

FIGURE 32 illustrates, in tabular format, T cell responses induced using a heterologous Ad34 prime/Ad35 boost regimen in macaques. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

FIGURE 33 illustrates, in tabular format, the levels of CD4+ and CD8+ Gagspecific T cells in Ad34 primed/Ad35 boosted macaques at week 28. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

## DETAILED DESCRIPTION OF THE INVENTION

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The present invention details an efficient strategy for the propagation and rescue of alternative adenoviral serotypes utilizing available adenovirus production cell lines, nullifying the need to customize available cell lines for a specific serotype of interest. This is enabled by the incorporation of a critical E4 region into the adenovirus to be propagated.

The critical E4 region in the instant invention comprises, in the minimum, nucleic acid sequence encoding E4 ORF6 and can comprise the entire region of E4, inclusive of the promoter region. An important characteristic of the imported E4 region is that it is native to a virus of the same or highly similar serotype as the E1 gene product(s) (particularly E1B 55K) of the E1-complementing cell line, but heterologous to (i.e., non-native to a virus of the same serotype as) the adenoviral vector being propagated. As will be detailed below, the heterologous E4 region or portion thereof can be varied and can be inserted into the vector backbone at numerous locations.

The heterologous E4 region or portion thereof can, for instance, be a nucleic acid sequence encoding the entire open reading frame of the non-native E4. This segment of nucleic acid sequence can, in turn, be incorporated into the "native" entire E4 open reading frame of the recipient virus. In such an embodiment, the promoter native to the adenoviral vector would drive the expression of the non-native E4 region within the recombinant replication-defective adenoviral vector. Alternatively, the nucleic acid sequence encoding the entire open reading frame can be inserted into a different region of the adenoviral vector genome, such as for example the E1 or E3 regions. In this latter embodiment, the native E4 region or portion thereof can be deleted or left intact.

In another embodiment, the heterologous E4 region comprises a nucleic acid sequence encoding the entire open reading frame of E4 and includes a non-native E4 promoter. In this type of embodiment, the E4 region can be inserted into the location of the combined native E4 and E4 promoter region. The non-native E4 region in this embodiment would be driven by expression of the non-native E4 promoter. Alternatively, the nucleic acid sequence encoding the entire open reading frame and the non-native E4 promoter can be inserted into a different region of the adenoviral vector genome, such as for example the E1 or E3 regions. In this latter embodiment, the native E4 region or portion thereof can be deleted or left intact.

An alternative and further embodiment exists wherein the heterologous E4 region or portion thereof comprises nucleic acid sequence encoding a partial E4 region comprising ORF6 (one aspect of which is a region solely encoding ORF6). In this particular aspect of the invention, the heterologous non-native E4 protein can, in certain embodiments, replace the non-native ORF6 region or the entire E4-encoding region of the native virus. In the latter situation, the promoter driving expression of the non-native ORF6 can either be the native E4 promoter or a heterologous, non-native promoter operatively linked to the non-native ORF6, while in the latter, the expression of the non-native ORF6 would generally be driven by the native E4 promoter. Alternatively, the nucleic acid sequence encoding a partial E4 region comprising ORF 6 can be inserted into a different region of the adenoviral vector genome, such as for example the E1 or E3 regions. In this latter embodiment, the native E4 region or portion thereof can be deleted or left intact.

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As one of skill in the art can appreciate, there are various ways in which one can envision the supply of a heterologous E4 nucleic acid sequence in cis to an adenoviral vector and thereby enable its growth based on Applicants' novel findings herein. Moreover, as one of skill in the art can appreciate, either native or non-native promoters can be utilized to drive expression of the heterologous E4 region or portion thereof.

Adenovirus pre-plasmids (plasmids comprising the genome of the replication-defective adenovirus with desired deletions and insertions) can be generated by homologous recombination using adenovirus backbones and an appropriate shuttle vector (designed to target-in specific deletions and incorporate desired restriction sites into the resultant plasmid). Shuttle vectors of use in this process can be generated using general methods widely understood and appreciated in the art, e.g., PCR of the adenoviral terminal ends taking into account the desired deletions, and the sequential cloning of the respective segments into an appropriate cloning plasmid. The adenoviral pre-plasmid can then be digested and transfected into the complementing cell line via calcium phosphate co-precipitation or other suitable means. Virus replication and amplification then occurs, a phenomenon made evident by notable cytopathic effect. Infected cells and media are then harvested after viral replication is complete (generally, 7-10 days post-transfection).

It is to be noted that various alternative adenoviral serotypes can be developed in accordance with the disclosed methods and, particularly, alternative adenoviral serotype vectors that were previously unable to be propagated or very inefficiently propagated utilizing existing adenoviral production cell lines based on subgroup C complementing E1 sequence. The various adenoviral vectors that can be developed in accordance with the instant methods include adenoviral vectors of subgroups A-F (for instance, serotypes of subgroups A, B (e.g., serotypes

11, 14, 16, 21, 34 and 35), C (e.g., serotypes 2 and 5), D (e.g., serotypes 24, 26 and 36), E (e.g., serotype 4) and F.

In preferred embodiments, the various non-group C family members can be developed with heterologous E4 supplied from a subgroup C member such as adenovirus serotype 5. Particular embodiments of the instant invention utilize a development scheme wherein the heterologous E4 protein is derived from a wildtype adenovirus serotype 5 sequence; see, e.g., a viral sequence which has been deposited with the American Type Culture Collection ("ATCC") under ATCC Deposit No. VR-5 (for which a transcription map can be found in Figure 1). A particular example of this type of embodiment is wherein an adenovirus of subgroup B (or any non-C subgroup) comprising heterologous E4 proteins in cis from Ad5 is propagated in Ad5 E1-complementing cell lines, for instance, PER.C6<sup>TM</sup> or 293. Applicants have, in fact, successfully propagated E1- serotypes 10, 24, 34, and 35 via use of this particular embodiment.

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One of skill in the art can readily identify alternative adenovirus serotypes (e.g., alternative serotypes of subgroups A, B (e.g., serotypes 11, 14, 16, 21, 34 and 35), C, (e.g., serotypes 2 and 5), D (e.g., serotypes 24, 26 and 36), E (e.g., serotype 4) and F) for the supply of the heterologous E4 protein. As long as the heterologous E4 region (or portion thereof comprising ORF6) of the vector is native to a virus of the same or highly similar serotype as the E1 region of the complementing cell line, the methods of the instant invention are widely applicable to the propagation and rescue of adenovirus of all serotypes. In light of the present disclosure, one can readily envision, for instance, how a complementing cell line based on a non-subgroup C adenovirus (e.g., the Ad35 cell line of EP 1 054 064 A1) can be utilized to propagate a virus of an adenoviral vector of subgroup C (e.g., adenovirus serotype 5) provided that the appropriate nucleic acid sequence encoding an E4 protein provided in cis is native to a virus of the same or highly similar serotype as that of the E1 expressed by the complementing cell line (i.e., an Ad35 E4 protein).

Complementing cell lines of use in the instant invention are available in the art and are not limited to any specific type. The critical feature, again, is that the heterologous segment of E4-encoding nucleic acid sequence provided in cis to the replication-defective vector being propagated be native to a virus of the same or highly similar serotype as the E1 expressed by the complementing cell line. One aspect of the instant invention employs E1-complementing cell lines wherein the expressed E1 is of serotype 5; e.g., PER.C6<sup>TM</sup> and 293 cell lines. Both these cell lines express the adenoviral E1 gene product. PER.C6<sup>TM</sup> is described in Fallaux et al., 1998 Human Gene Therapy 9:1909-1917, hereby incorporated by reference. 293 cell lines are described in Graham et al., 1977 J. Gen. Virol. 36:59-72, hereby incorporated by reference.

Another aspect of the instant invention are the adenoviral vectors of any serotype falling with adenoviral subgroups A, B, C, D, E and F (for instance, alternative serotypes of subgroups A, B (e.g., serotypes 11, 14, 16, 21, 34 and 35), C (e.g., serotype 2), D (e.g., serotypes 24, 26 and 36), E (e.g., serotype 4) and F) which are modified to contain a non-native E4-encoding nucleic acid sequence in cis which comprises, in whole or in part, nucleic acid sequence encoding open reading frame 6 (ORF6). Virus in accordance with this description can be propagated in accordance with the above-described methods and rescued using any suitable means known in the art.

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Another aspect of the instant invention is a vector in accordance with the instant invention which comprises a heterologous passenger gene in addition to that of the heterologous E4 nucleic acid sequence. In specific embodiments, the passenger gene encodes an antigen.

As one of ordinary skill in the art will appreciate, the instant methods are not limited by the heterologous gene that can be incorporated. The instant invention relates generally to a means by which to propagate multiple serotypes of adenovirus in a single complementing cell line and the recombinant virus that make the process possible. In preferred embodiments, the passenger gene is incorporated into the E1 deletion. In alternatively preferred embodiments, the passenger gene is inserted in an E3-deleted region. The position of the passenger gene, as one of ordinary skill in the art will appreciate, can be varied according to the specific complementing cell utilized and the specific deletions present within the replication-defective adenovirus genome.

In specific embodiments the passenger gene can encode an HIV-1 antigen, and in more preferred embodiments selected from the group consisting of genes encoding HIV-1 gag, pol, nef and env. In alternative embodiments, the passenger gene can be a reporter gene, such as secreted alkaline phosphatase (SEAP).

The passenger gene preferably exists in the form of an expression cassette. A gene expression cassette preferably comprises (a) a nucleic acid sequence encoding a protein of interest; (b) a promoter operatively linked to the nucleic acid sequence encoding the protein; and (c) a transcription termination sequence. The transcriptional promoter of the adenoviral vector is preferably recognized by an eukaryotic RNA polymerase. In a preferred embodiment, the promoter is a "strong" or "efficient" promoter. An example of a strong promoter is the immediate early human cytomegalovirus promoter (Chapman et al., 1991 Nucl. Acids Res. 19:3979-3986), which is hereby incorporated by reference), in certain embodiments without intronic sequences. Those skilled in the art, however, will appreciate that any of a number of other known promoters, such as the strong immunoglobulin, or other eukaryotic gene promoters

may also be used, including the EF1 alpha promoter, the murine CMV promoter, Rous sarcoma virus (RSV) promoter, SV40 early/late promoters and the beta-actin promoter.

The promoter may comprise a regulatable sequence such as the Tet operator sequence. This is extremely useful, for example, in cases where the gene products are affecting a result other than that desired and repression is sought.

Transcription termination sequences can also be utilized within the gene expression cassettes. Preferred termination sequences are, for instance, the bovine growth hormone terminator/polyadenylation signal (bGHpA) and the short synthetic polyA signal (SPA) of 50 nucleotides in length, defined as follows:

Further embodiments incorporate a leader or signal peptide into the transgene. A preferred leader is that from the tissue-specific plasminogen activator protein, tPA.

The following non-limiting Examples are presented to better illustrate the

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#### EXAMPLE 1

#### Construction and Rescue

An E1- Ad35-based pre-adenovirus plasmid was constructed in order to determine whether an E1- Ad35 vector (a representative group B serotype) could be propagated in a group C E1-complementing cell line. The general strategy used to recover Ad35 as a bacterial plasmid is illustrated in Figure 3. Cotransformation of BJ5183 bacteria with purified wild-type Ad35 viral DNA and a second DNA fragment termed the Ad35 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The ITR cassette contains sequences from the right (bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome (see Figures 2A-1 to 2A-10) separated by plasmid sequences containing a bacterial origin of replication and an Ampicillin resistance gene. The ITR cassette contains a deletion of E1 sequences from Ad5 457 to 3402 with a unique Swa I site located in the deletion. The Ad35 sequences in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which recombination can occur. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1. Pre-Adenovirus plasmid pAd35ΔE1 contains Ad35 sequences from 4 to 456 and bp 3403 to 34793.

To determine if pre-adenovirus plasmid pAd35ΔE1 could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into a T-25 flask of PER.C6 cells using the calcium phosphate co-precipitation technique. *Pme* I digestion releases the viral genome from the plasmid sequences allowing viral replication to occur after entry into 293 cells. Viral cytopathic effect (CPE), indicating that virus replication and amplification is occurring, was never observed. Cells and media from the transfection were harvested at 14 days post transfection, freeze-thawed three times, clarified by centrifugation and used to infect new PER.C6 cells but no virus was ever amplified. Following multiple attempts, we have been unable to rescue and amplify pAd35ΔE1 in PER.C6 cells.

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#### **EXAMPLE 2**

## Insertion of Ad5 Orf 6 and Ad5 E4 into the Ad5 Genome

To refine the strategy of including Ad5 Orf6 in the genome of an alternative serotype so that propagation could take place in a Ad5/group C complementing cell line four additional strategies were developed. In the first strategy, the entire alternative serotype E4 region (not including the E4 promoter) was deleted and replaced with Ad5 Orf6. In the second strategy, just the alternative serotype Orf6 gene was deleted and replaced with Ad5 Orf6. In the third strategy, the entire alternative serotype E4 coding region (not including the E4 promoter) was deleted and replaced with the Ad5 E4 coding region (not including the Ad5 E4 promoter) and, in the final strategy, the entire alternative serotype E4 coding and promoter region was deleted and replaced with the Ad5 E4 promoter and coding region. The configuration of the E4 regions generated by the four strategies is diagramed in Figure 4. For each of these strategies the desired pre-Adenovirus plasmid was generated by bacterial recombination. Cotransformation of BJ 5183 bacteria with purified wild-type viral DNA and the appropriately constructed ITR cassette resulted in the circularization of the viral genome by homologous recombination. The construction of each pre-Ad plasmid, based on Ad35, is outlined below:

To construct pAd35 $\Delta$ E1 $\Delta$ E4Ad5Orf6 (An Ad35 pre-Ad plasmid containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31913 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 31913 and 34419 generating pNEBAd35-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a

bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31912 to 34418 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct, Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31913 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria (Figure 5). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔE4Ad5Orf6. Pre-Adenovirus plasmid pAd35ΔE1ΔE4Ad5Orf6 contains Ad35 sequences from bp 4 to 456; bp 3403 to bp 31913 and bp 34419 to bp 34793 with Ad5Orf6 cloned between bp 31913 and bp 34419.

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To construct pAd35\Delta E1\Delta Orf6Ad5Orf6 (An Ad35 pre-Ad plasmid containing an E1 deletion and a deletion of E4 Orf6 substituted with Ad5 Orf6), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 32081 and bp 32990 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-10. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 32081 and 32990 generating pNEBAd35-10Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and a deletion of E4 Orf6 from Ad35 bp 32082 to 32989 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct, Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 32081 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔOrf6Ad5Orf6. Pre-Adenovirus plasmid pAd35ΔE1ΔOrf6Ad5Orf6 contains Ad35

sequences from bp 4 to 456; bp 3403 to bp 32081 and bp 32990 to bp 34793 with Ad5Orf6 cloned between bp 32081 and bp 32990.

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To construct pAd35ΔE1ΔE4Ad5E4 (An Ad35 pre-Ad plasmid containing an E1 deletion and a deletion of E4 substituted with Ad5 E4), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31838 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-7. Next the Ad5 E4 coding region was generated by PCR and cloned between Ad35 bp 31838 and 34419 generating pNEBAd35-7Ad5E4-2 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31839 to 34418 into which the Ad5 E4 coding region was introduced in an E4 parallel orientation. In this construct, the Ad5 E4 region is expressed using the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31838 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔE4Ad5E4. Pre-Adenovirus plasmid pAd35ΔE1ΔE4Ad5E4 contains Ad35 sequences from bp 4 to 456; bp 3403 to bp 31838 and bp 34419 to bp 34793 with the Ad5 E4 coding region (Ad 5 bp 32914 to bp 35523) cloned between bp 31838 and bp 34419.

To construct pAd35\DelaE1\DelaE4Ad5PE4 (An Ad35 pre-Ad plasmid containing an E1 deletion and a deletion of E4 coding region and promoter substituted with Ad5 E4 coding region and promoter), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31838 and bp 34660 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-8. Next the Ad5 E4 promoter and coding region was generated by PCR and cloned between Ad35 bp 31838 and 34660 generating pNEBAd35-8Ad5E4PC (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication,

ampicillin resistance gene, and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31839 to 34659 into which the Ad5 E4 promoter and coding region was introduced in an E4 parallel orientation. In this construct, the Ad5 E4 region is expressed using the Ad5 E4 promoter. The Ad35 sequences (bp 31599 to 31838 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔE4Ad5PE4. Pre-Adenovirus plasmid pAd35ΔE1ΔE4Ad5PE4 contains Ad35 sequences from bp 4 to 456; bp 3403 to bp 31838 and bp 34660 to bp 34793 with the Ad5 E4 promoter and coding region (Ad 5 bp 32914 to bp 35826) cloned between bp 31838 and bp 34660.

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#### EXAMPLE 3

Rescue of pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4 into Virus

In order to determine if pre-adenovirus plasmids pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4 could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmids were each digested with Pme I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique; Cell Phect Transfection Kit, Amersham Pharmacia Biotech Inc. PmeI digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring, was observed for all construct. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Approximately 1 ml of the cell lysate was used to infect aT-225 flasks of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following complete CPE the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then

digested with *Hind*III and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with *Pme1/Hind*III prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

#### **EXAMPLE 4**

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Insertion of an Expression Cassette into pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4

In order to introduce a gag or SEAP expression cassette into the E1 region of the various Ad35 pre-Adenovirus plasmids described above (pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4) bacterial recombination was again used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence (Figure 6), was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2 (a precursor to the Ad35 ITR cassettes described above), generating pNEBAd35CMVgagBGHpA. pNEBAd35-2 contains Ad35 sequences from the left end of the genome (bp 4 to 456 and bp 3403 to 3886) with a unique Swal site between bp 456 and 3403 at the position of the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pNEBAd35-2. This cloning step resulted in the gag expression cassette being cloned into the E1 deletion between bp 456 and 3403 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad35 bp 4 to 456 and bp 3403 to 3886 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and one of the Ad35 pre-Ad plasmids (pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4, pAd35ΔE1ΔE4Ad5PE4), linearized in the E1 region by digestion with Swa I, resulted in the generation of corresponding Ad35 gag-containing pre-Adenovirus plasmids (pAd35ΔE1gagΔE4Ad5Orf6, pAd35ΔE1gagΔOrf6Ad5Orf6, pAd35 $\Delta$ E1gag $\Delta$ E4Ad5E4, and pAd35 $\Delta$ E1gag $\Delta$ E4Ad5PE4) by homologous recombination. Potential clones were screened by restriction analysis.

A similar strategy was used to generate Ad35 pre-Ad plasmids containing a SEAP expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence (Figure 7) was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2, generating pNEBAd35CMVSEAPBGHpA. The SEAP expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pNEBAd35-2. The transgene was then recombined into the various Ad35 backbones generating pAd35ΔE1SEAPΔE4Ad5Orf6, pAd35ΔE1SEAPΔC4Ad5Orf6, pAd35ΔE1SEAPΔE4Ad5E4, and pAd35ΔE1SEAPΔE4Ad5PE4 as described above for the gag transgene. All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

#### 15 EXAMPLE 5

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In vivo Transgene Expression

#### A. Immunization

Female mice were between 4-10 weeks old. The total dose of each vaccine was suspended in 0.1 mL of buffer. The vectors were given to both quadriceps of each animals with a volume of 50 µL per quad and using 0.3-mL 28G1/2 insulin syringes (Becton-Dickinson, Franklin Lakes, NJ). The rhesus macaques and African green monkeys were between 2-5 kg in weight. For the primates, the total dose of each vaccine was suspended in 1 mL of buffer. The monkeys were anesthetized (ketamine/xylazine mixture) and the vaccines were delivered i.m. in 0.5-mL aliquots into two muscle sites using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Serum samples were collected at defined intervals and stored frozen until the assay date. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council.

#### B. SEAP Assay

Serum samples were analyzed for circulating SEAP levels using TROPIX phospha-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5  $\mu$ L aliquots of each serum were mixed with 45  $\mu$ L of kit-supplied dilution buffer in a 96-well white DYNEX plate.

Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey or mouse serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the well for 30 minutes at 65 °C. Enzymatic SEAP activities in the samples were determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorder using DYNEX luminometer. RLU readings are converted to ng/mL SEAP using a log-log regression analyses.

#### C. Rodent Results

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In the first mouse experiment, cohorts of 5 C3H/HeN mice were given single intramuscular injections of one of the following vectors: (1) 10^10 vp Ad35ΔE1SEAPΔE4Ad5Orf6; (2) 10^10 vp Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6; or (3) 10^10 vp Ad35ΔE1SEAP. Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 8. Results indicate that (1) the Ad35 constructs are all capable of expressing the SEAP transgene and that (2) the introduction of Ad5Orf6 sequence where the deleted Ad35E4 was did not significantly affect the transgene expression relative to Ad35ΔE1SEAP. Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6 also yielded a similar expression profile as Ad35ΔE1SEAP. The levels of SEAP in the serum dropped after day 2 and were at background levels by day 12.

The second mouse experiment evaluates the effect of a full Ad5E4 replacement instead of an Ad5Orf6 substitution for the Ad35 E4 cassette. Here, cohorts of 5 C3H/HeN mice were given single intramuscular injections of one of the following vectors: (1) 10^10 vp MRKAd5-SEAP; (2) 10^9 vp MRKAd5-SEAP; (3) 10^10 vp Ad35ΔE1SEAPΔE4Ad5Orf6; (4) 10^10 vp Ad35ΔE1SEAPΔE4Ad5E4; or (5) 10^10 vp Ad35ΔE1SEAPΔE4Ad5PE4. The introduction of Ad5E4 or Ad5PE4 resulted in comparable if not, slightly improved expression levels compared to the vector with the Ad5Orf6 insertion (Figure 9). The peak levels for the Ad35 constructs are lower than those produced by Ad5SEAP (at least 10-fold).

#### D. Primate Results

Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the following vectors: (1) 10^11 vp MRKAd5-SEAP; (2) 10^9 vp MRKAd5-SEAP; or (3) 10^11 vp Ad35\Delta E1SEAP\Delta E4Ad5Orf6. Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results for the individual monkeys are shown in Figures 10A-B. Results indicate that the peak level of SEAP product produced by the alternative adenovirus serotype was lower than but were within 3-fold of that of MRKAd5SEAP at the same

high dose level of 10^11 vp. The levels observed from the Ad35 vector were about 50-fold higher than those observed using 10^9 vp of MRKAd5SEAP. The levels of SEAP in the serum dropped after day 10 and were close to background as early as day 15.

A separate experiment using African green monkeys was conducted to examine the effect of the additional E3 deletion or the full Ad5E4 substitution on in vivo gene expression. In here, cohorts of 2-3 African green macaques were given single intramuscular injections of one of the following vectors: (1) 10^11 vp MRKAd5-SEAP; (2) 10^10 vp MRKAd5-SEAP; (3) 10^9 vp MRKAd5-SEAP; (4) 10^10 vp Ad35ΔE1SEAPΔE4Ad5Orf6; (5) 10^10 vp Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6; or (6) 10^10 vp Ad35ΔE1SEAPΔE4Ad5E4. Results (Figure 11) indicate that the peak levels of SEAP product produced by Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6 and Ad35ΔE1SEAPΔE4Ad5E4 were comparable if not, slightly improved compared to Ad35ΔE1SEAPΔE4Ad5Orf6.

#### **EXAMPLE 6**

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In vivo Immunogenicity

#### A. Immunization

Cohorts of 3-6 animals were given intramuscular injections at wk 0 and wk 4 of either of the following constructs: (1) 10^11 vp MRKAd5-HIV1 gag; or (2) 10^11 vp of Ad35ΔE1gagΔE4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson). Sera and peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the *Guide for Care and Use of Laboratory Animals*, Institute of Laboratory Animal Resources, National Research Council.

#### 30 B. ELISPOT Assay

The IFN-γ ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen *et al.*, 2001 *J. Virol.* 75(2):738-749), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50 μL of 2-4 x 10<sup>5</sup> peripheral blood mononuclear cells (PBMCs)

were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower size cut-off set at 80 femtoliters ("fL"). Either 50 μL of media or the gag peptide pool at 8 μg/mL concentration per peptide was added to the PBMC. The samples were incubated at 37°C, 5% CO<sub>2</sub> for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform (Silver Spring, MD); the counts were normalized to 10<sup>6</sup> cell input.

## C. Intracellular Cytokine Staining

To 1 ml of 2 x 10<sup>6</sup> PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added to a final concentration of 1  $\mu$ g/mL. For gag-specific stimulation, 10  $\mu$ L of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20  $\mu L$  of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO<sub>2</sub>, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20 µL per tube antihCD3-APC, clone FN-18 (Biosource); 20 μL anti-hCD8-PerCP, clone SK1 (Becton Dickinson, Franklin Lakes, NJ); and 20 µL anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750  $\mu L$ 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in PBS/2%FBS and 0.1 μg of FITC-anti-hIFN-γ, clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence cut-off for cytokine-positive events was used for both CD4+ and CD8+ populations, and for both mock and gag-peptide reaction tubes of a sample.

#### 30 D. Results

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PBMCs collected at regular 4-wk intervals were analyzed in an ELISPOT assay. Results (Table 1) indicate that the Ad35ΔE1gagΔE4Ad5Orf6 is able to induce in non-human primates significant levels of gag-specific T cells. After a single dose (wk 4), the Ad35-induced responses were about 5-fold lower than that of MRKAd5-HIV1 gag. After the second dose (wk

8), the responses between both cohorts were comparable; the differences became pronounced in the succeeding time points.

Table 1. Gag-specific T cell response in monkeys immunized with MRKAd5-HIV1gag and Ad35ΔE1gagΔE4Ad5Orf6. Shown is the number of spot-forming cells per million PBMC following incubation in the absence (mock) or presence of Gag H peptide pool. The H pool consisted of 20-aa peptide overlapping by 10 aa and encompassing the entire gag sequence.

Grp	Vaccine Wk 0, Wk 4	Monkey ID	Pre		Wk 4		Wk 8		Wk 12		Wk 16	
Gib			Mock	Gag H	Mock	Gag H	Mock	Gag H	Mock	Gag H	Mock	Gag H
1	MRKAd5-HIV1 gag 10^11 vp	00C018 00C034 00C058	1 0 4	5 4 4	13 5 3	1025 219 1086	0 5 0	824 404 440	3 0 0	753 491 439	1 1 0	533 350 599
2	Ad35aE1gagAE4Ad5Orf6 10^11 vp	00D045 00D067 00D068 00D054 00D075 00D073	1 1 1 3 3	1 4 4 15 5 26	3 5 10 10 18 1	168 89 34 195 275 241	5 0 5 0 13 3	645 103 365 501 716 485	4 0 3 3 3	178 76 143 350 158 278	0 0 0 0	91 19 95 124 103 148
3	Naīve	00D087	1	1	3	3	8	54	3	5	3_	1 1

Intracellular IFN-γ staining analyses of PBMC collected at wk 8 suggest that the Ad35-based vaccine is able to induce both HIV-specific CD4+ and CD8+ T cells (Table 2).

Table 2. Characterization of the gag-specific T cells in monkeys immunized with MRKAd5-HIV1gag and Ad35ΔE1gagΔE4Ad5Orf6. Shown are the percentages of CD3+ T cells that are either gag-specific CD4+ or gag-specific CD8+ cells. These values were corrected for mock values (<0.02%).

Grp	Vaccine Wk 0, Wk 4	Monkey	Wk 8			
		ID	%CD4+CD3+	%CD8+CD3+		
1	MRKAd5-HIV1 gag	00C018	0.08	0.37		
•	10^11 vp	00C034	0.09	0.06		
	•	00C058	0.03	0.21		
2	Ad35AE1gagAE4Ad5Orf6	00D045	0.06	0.08		
_	10^11 vp	00D067	0.02	0.02		
		00D068	0.15	0.02		
	•	00D054	0.05	0.08		
		00D075	0.08	0.05		
		00D073	0.09	0.06		

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In a separate experiment, 3 different Ad35 constructs expressing HIV-1 gag were evaluated for their immunogenicity in macaques. Here, cohorts of 3 macaques were given immunizations at wk 0 and 4 of either of the following vectors: (1) 10^10 vp Ad35ΔE1gagΔE4Ad5Orf6; (2) 10^10

vp Ad35ΔE1gagΔE3ΔE4Ad5Orf6; or (3) 10^10 vp Ad35ΔE1gagΔE4Ad5E4. The levels of T cell immunity induced by all 3 vectors were comparable at this stage (Table 2), suggesting that the additional E3 deletion or full Ad5E4 substitution does not appear to impair the immunogenic properties of the vector.

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Table 3. Gag-specific T cell response in monkeys immunized with several Ad35ΔE1ΔE4-based vectors. Shown is the number of spot-forming cells per million PBMC following incubation in the absence (mocK0 or presence of Gag H peptide pool. The H pool consisted of 20-aa peptide overlapping by 10 aa and encompassing the entire gag sequence.

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Grp	Vaccine	Monkey	Pre		Wk 4		Wk 8	
Gib	Wk 0, Wk 4	ID	Mock	Gag H	Mock	Gag H	Mock	Gag H
	Ad35∆E1gag∆E4Ad5Orf6	00C047	4	1	0	20	0	189
'	10^10vp	00C157	8	5	1	81	1	833
.	10 104β	00C078	3	1	0	46	4	349
2	Ad35∆E1gag∆E3∆E4Ad5Orf6	00C091	1	1	1	118	3	315
2	10^10vp	00C122	3	lo	0	31	· 1	138
	ιστιονρ	00D177	3	3	1	45	1	64
3	Ad35∆E1gag∆E4Ad5E4	00D018	3	19	29	120	23	193
	10^10vp	00D046	8	5	1	21	10	143
	10 1000	00D063	3	4	0	63	4	371
Naïve	none	00D363	0	5	ND	ND	0	0
IVAIVE							L	

### EXAMPLE 7

## Construction and Rescue of pAd24ΔE1.

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An E1- Ad24-based pre-adenovirus plasmid was constructed in order to determine whether an E1- Ad24 vector (a representative group D serotype) could be propagated in an Ad5/group C E1-complementing cell line. Since at the time the vector construction was initiated the complete sequence of Ad24 (see Figures 16A-1 through 16A-10; subject of copending application serial no. 60/455, 312, filed March 17, 2003) was unknown we took advantage of some sequence homology between Ad24 and Ad17. The general strategy used to recover Ad24 as a bacterial plasmid is illustrated in Figure 12 and described below.

Cotransformation of BJ5183 bacteria with purified wild-type Ad24 viral DNA and a second DNA fragment termed the Ad17 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The ITR cassette contains sequences from the right (bp 34469 to 35098) and left (bp 4 to 414 and bp 3373 to 4580) end of the Ad17 genome (Accession No. AF108105) separated by plasmid sequences containing a bacterial origin of replication and an Ampicillin resistance gene. The ITR cassette contains a deletion of E1 sequences from Ad17

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(bp 415 to 3372) with a unique Swa I site located in the deletion. The Ad17 sequences in the ITR cassette provide regions of homology with the purified Ad24 viral DNA in which recombination can occur. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the Ad24 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd24ΔE1. pAd24ΔE1 contains Ad17 sequences from bp 4 to 414 and from bp 3373 to 4580, Ad24 bp 4588 to 34529, and Ad17 bp 34469 to 35098 (bp numbers refer to the wt sequence for both Ad17 and Ad24). PAd24ΔE1 contains the coding sequences for all Ad24 virion structural proteins that constitute its serotype specificity. This approach can be used to circularize any group D serotype into plasmid form which has sufficient homology to Ad17.

To determine if pre-adenovirus plasmid pAd24ΔE1 could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into a 6 cm dish of 293 cells using the calcium phosphate co-precipitation technique. *Pme* I digestion releases the viral genome from the plasmid sequences allowing viral replication to occur after entry into 293 cells. Viral cytopathic effect (CPE), indicating that virus replication and amplification is occurring, was very slow to arise. Following multiple attempts, we were successful at rescuing and amplifying Ad24ΔE1 but the virus grew to lower titers and took more passages to amplify than a similar Ad5 based vector. In order to verify the genetic structure of the virus, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with *Hind*III and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products from the pre-plasmid (that had been digested with *Pme1/Hind*III prior to labeling). The expected sizes were observed, indicating that the virus had been successfully rescued.

#### **EXAMPLE 8**

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### Insertion of Ad5 Orf 6 into the E1 region of Ad24

In order to determine if the insertion of Ad5 E4 Orf6 into the Ad24 genome would allow more efficient propagation in a group C E1 complementing cell line we constructed an Ad24 based pre-adenovirus plasmid containing Ad5 Orf6 in the E1 region. In order to introduce Ad5 Orf6 in to the E1 region of pAd24ΔE1, bacterial recombination was used. An Ad5 Orf6 transgene consisting of the Ad5 Orf6 coding region flanked by the HCMV promoter and pA was cloned into the E1 deletion in an Ad17 shuttle vector (a precursor to the Ad17 ITR cassette). The Ad5 Orf6 transgene was cloned between bp 414 and 3373 in the E1 anti-parallel

orientation. The shuttle vector containing the Ad5 Orf6 transgene was digested to generate a DNA fragment consisting of the transgene flanked by Ad17 sequences (bp 4 to 414 and bp 3373 to 4580) and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pAd24ΔE1, which had been linearized in the E1 region by digestion with *Swa*I, resulted in the generation of pAd24ΔE1Ad5Orf6 by homologous recombination (Figure 13). Potential clones were screened by restriction analysis and one clone was selected as pre-adenovirus plasmid pAd24ΔE1Ad5Orf6.

In order to determine if pre-adenovirus plasmid pAd24ΔE1Ad5Orf6 could be rescued into virus and propagated in an Ad5/group C E1 complementing cell line, pAd24ΔE1Ad5Orf6 was digested with Pme I and transfected into a 6 cm dish of 293 cells using the calcium phosphate co-precipitation technique. PmeI digestion releases the viral genome from plasmid sequences allowing viral replication to occur after entry into 293 cells. Once complete viral cytopathic effect (CPE) was observed at approximately 7-10 days post transfection, the infected cells and media were freeze/thawed three times and the cell debris pelleted. The virus was amplified in two additional passages in 293 cells and then purified from the final infection by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the virus, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with HindIII and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The endlabeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products from the pre-plasmid (that had been digested with Pme1/HindIII prior to labeling). The expected sizes were observed, indicating that the virus had been successfully rescued.

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### **EXAMPLE 9**

## Insertion of Ad5 Orf 6 into the E4 region of Ad24

To refine the strategy of including Ad5 Orf6 in the genome of an alternative serotype so that propagation could take place in an Ad5/group C complementing cell line two additional strategies were developed. In the first strategy, the entire alternative serotype E4 region (not including the E4 promoter) was deleted and replaced with Ad5 Orf6. In the second strategy, just the alternative serotype Orf6 gene was deleted and replaced with Ad5 Orf6. The configuration of the E4 regions generated by the two strategies is diagramed in Figure 14. For each of these strategies the desired pre-Adenovirus plasmid was generated by bacterial recombination. Cotransformation of BJ 5183 bacteria with pAd24ΔOrf6BstZ17I and the

appropriately constructed Ad24 E4 shuttle plasmid resulted in the generation of the desired Ad24 based pre-Ad plasmid. PAd24ΔOrf6BstZ17I, a derivative of pAd24ΔE1, was constructed so that the E4 region in the Ad24 pre-Ad plasmid could be easily modified using bacterial recombination. PAd24ΔOrf6BstZ17I contains a deletion in the E4 region from Ad24 bp 32373 to bp 33328 with a unique BstZ17I site located at the position of the deletion. The complete sequence of pAd24ΔOrf6BstZ17I consists of Ad17 sequences from bp 4 to 414 and from bp 3373 to 4580, Ad24 bp 4588 to 32372 and from 33329 to 34529, and Ad17 bp 34469 to 35098 (bp numbers refer to the wt sequence for both Ad17 and Ad24).

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To construct pAd24ΔE1ΔE4Ad5Orf6 (An Ad24 pre-Ad plasmid containing an E1 deletion and a deletion of E4 substituted with Ad5 Orf6), an Ad24 E4 shuttle plasmid was constructed by digesting pAd24ΔE1 with *PmeI* and *BsrGI* and cloning the restriction fragment representing the E4 region (bp 31559 to bp 35164) into pNEB193, generating pNEBAd24E4. PNEBAd24E4 was then digested with *AccI* and *Eco*NI to remove the E4 coding sequences and ligated with an oligo designed to contain *BgIII* and *XhoI* sites (underlined) (5'

15 ACTCGAGATGTATAGATCT (SEQ ID NO: 6); 5' CTAGATCTATACATCTCGAG (SEQ ID NO: 7)), generating pNEBAd24ΔE4. PNEBAd24ΔE4 was then digested with BgIII and XhoI and ligated with the Ad5 Orf6 gene, which was PCR amplified, generating pNEBAd24ΔE4Ad5Orf6. The PCR primers used to amplify the Ad5 Orf6 gene (5' GCACAGATCTTTGCTTCAGGAATATG (SEQ ID NO: 8); 5'

GAGAACTCGAGGCCTACATGGGGGTAGAG (SEQ ID NO: 9)) were designed to contain BgIII and XhoI sites (underlined above) for ligation with the pNEBAd24DE4 fragment. In the final step pNEBAd24ΔE4Ad5Orf6 E4 shuttle plasmid was digested with PvuI and PmeI, the restriction fragments were size fractionated by agarose gel electrophoresis and the desired fragment containing Ad5Orf6 flanked by Ad24 sequences was gel purified. Cotransformation of BJ 5183 bacteria with E4 shuttle fragment and pAd24ΔOrf6BstZ17I, which had been linearized in the E4 region by digestion with BstZ17I, resulted in the generation of pAd24ΔE1ΔE4Ad5Orf6 by homologous recombination. Potential clones were screened by restriction analysis and one clone was selected as pre-adenovirus plasmid pAd24ΔE1ΔE4Ad5Orf6.

To construct pAd24ΔE1ΔOrf6Ad5Orf6 (An Ad24 pre-Ad plasmid containing an E1 deletion and a deletion of E4 Orf6 substituted with Ad5 Orf6), an Ad24 E4 shuttle plasmid was constructed in which the Ad24 Orf6 gene was replaced by Ad5 Orf6. To do this the *EcoR1* restriction fragment representing bp 32126 to bp 33442 of the Ad24 genome (encompassing the E4 Orf6 coding region), was subcloned into the *EcoR1* site in pNEB193, generating pNEBAd24Orf6. In order to delete the E4 Orf6 gene in pNEBAd24Orf6 and replace it with Ad5 Orf6, pNEBAd24Orf6 was digested with *StyI* and treated with Klenow to blunt the ends and then

digested with to EagI. The desired pNEBAd24Orf6 fragment was then ligated with a PCR product representing the Ad5 Orf6 gene from Ad5 bp 33193 to bp 24125, generating pNEBAd24ΔOrf6Ad5Orf6. The PCR primers used to generate the Ad5 Orf6 fragment (5'CGAGACGCCGACGCAGATCTGTTTG (SEQ ID NO: 10);

5 'GAAGTCCCGGGCTACATGGGGGTAG (SEQ ID NO: 11)) were designed to contain EagI and SmaI sites (underlined above) for ligation with the pNEBAd24Orf6 fragment. In the final step pNEBAd24ΔOrf6Ad5Orf6 was digested with EcoRI, the restriction fragments were size fractionated by agarose gel electrophoresis and the desired fragment containing Ad5Orf6 flanked by Ad24 sequences was gel purified. Cotransformation of BJ 5183 bacteria with the EcoRI fragment and pAd24ΔOrf6BstZ17I, which had been linearized in the E4 region by digestion with BstZ17I, resulted in the generation of pAd24ΔE1ΔOrf6Ad5Orf6 by homologous recombination. Potential clones were screened by restriction analysis and one clone was selected as preadenovirus plasmid pAd24ΔE1ΔOrf6Ad5Orf6.

#### 15 EXAMPLE 10

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## Rescue of pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6, into Virus

In order to determine if pre-adenovirus plasmids pAd24AE1AE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6, could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmids were each digested with Pme I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique; (Cell Phect Transfection Kit, Amersham Pharmacia Biotech Inc.). PmeI digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring, was observed for both constructs. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Approximately 1 ml of the cell lysate was used to infect T-225 flasks of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following complete CPE the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with HindIII and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared

with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with *Pme1/Hind*III prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

#### 5 EXAMPLE 11

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## Comparison of the Growth Kinetics of Ad24 based vectors.

In order to compare the growth kinetic of Ad24ΔE1, Ad24ΔE1Ad5Orf6, Ad24ΔE1ΔE4Ad5Orf6 and Ad24ΔE1ΔOrf6Ad5Orf6 one step growth curves were preformed (Figure 15). PER.C6 cells in 60 mm dishes were infected at 1 vp per cell with either Ad24ΔE1, Ad24ΔE1Ad5Orf6, Ad24ΔE1ΔE4Ad5Orf6 or Ad24ΔE1ΔOrf6Ad5Orf6. Cells and media were then harvested at various times post infection, freeze thawed three times and clarified by centrifugation. The amount of virus present in the samples was determined by quantitative PCR and is illustrated in Figure 15. This study demonstrates that Ad24 vectors that incorporate Ad5 Orf6 have a distinct growth advantage over Ad24ΔE1 in PER.C6 cells. The instant invention can be practiced with recombinant Ad24 vectors absent a heterologous Orf 6 region where the E1-complementing cell line expresses an Ad24 E1 region or, alternatively, E1 and E4 regions of the same serotype (such as Ad5E1/E4-expressing cell lines).

#### EXAMPLE 12

Insertion of an Expression Cassette into pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6, In order to introduce a gag or SEAP expression cassette (see Figures 6 and 7, respectively) into the E1 region of the Ad24 pre-Adenovirus plasmids described above (pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6) bacterial recombination was used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence, was cloned into the E1 deletion in Ad17 shuttle plasmid, pABSAd17-3, generating pABSAd17HCMVgagBGHpA. The ITR cassette contains sequences from the right (bp 34469 to 35098) and left (bp 4 to 414 and bp 3373 to 4580) end of the Ad17 genome separated by plasmid sequences containing a bacterial origin of replication and an Ampicillin resistance gene. The ITR cassette contains a deletion of E1 sequences from Ad17 (bp 415 to 3372) with a unique Swa I site located in the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the Swal site in pABSAd17-3. This cloning step resulted in the gag expression cassette being

cloned into the E1 deletion between bp 414 and 3373 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad17 bp 4 to 414 and bp 3373 to 4580 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and one of the Ad24 pre-Ad plasmids (pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6,), linearized in the E1 region by digestion with Swa I, resulted in the generation of the corresponding Ad24 gag-containing pre-Adenovirus plasmids (pAd24ΔE1gagΔE4Ad5Orf6, pAd24ΔE1gagΔOrf6Ad5Orf6) by homologous recombination. Potential clones were screened by restriction analysis.

A similar strategy was used to generate Ad24 pre-Ad plasmids containing a SEAP expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence was cloned into the E1 deletion in Ad17 shuttle plasmid, pABSAd17-3, generating pABSAd17HCMVSEAPBGH. The SEAP expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pABSAd17-3. The shuttle vector containing the SEAP transgene was digested to generate a DNA fragment consisting of the SEAP expression cassette flanked by Ad17 bp 4 to 414 and bp 3373 to 4580 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and one of the Ad24 pre-Ad plasmids (pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6,), linearized in the E1 region by digestion with Swa I, resulted in the generation of the corresponding Ad24 SEAP-containing pre-Adenovirus plasmids (pAd24ΔE1SEAPΔE4Ad5Orf6, pAd24ΔE1SEAPΔOrf6Ad5Orf6) by homologous recombination. Potential clones were screened by restriction analysis. All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

#### **EXAMPLE 13**

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30 In Vivo Immunogenicity

#### A. Immunization

Cohorts of 3-6 animals were given intramuscular injections at wk 0 and wk 4 of either of the following constructs: (1) 10^11 vp MRKAd5-HIV1 gag; (2) 10^10 vp MRKAd5-HIV1 gag; (3) 10^11 vp of Ad24ΔE1gagΔOrf6Ad5Orf6; (4) 10^10 vp of

Ad24ΔE1gagΔOrf6Ad5Orf6; or (5) 10^10 vp of Ad24ΔE1gagΔE4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points (typically 4 wk intervals) during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the *Guide for Care and Use of Laboratory Animals*, Institute of Laboratory Animal Resources, National Research Council.

#### B. ELISPOT Assay

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The IFN-γ ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen et al., 2001 *J. Virol.* 75(2):738-749; Casimiro et al., 2002 *J. Virol.* 76:185-94), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50 μL of 2-4 x 10<sup>5</sup> peripheral blood mononuclear cells (PBMCs) were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower size cut-off set at 80 femtoliters ("fL"). Either 50 μL of media or the gag peptide pool at 8 μg/mL concentration per peptide was added to the PBMC. The samples were incubated at 37°C, 5% CO<sub>2</sub> for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform (Silver Spring, MD); the counts were normalized to 10<sup>6</sup> cell input.

#### C. Intracellular Cytokine Staining

To 1 ml of 2 x 10<sup>6</sup> PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added to a final concentration of 1 μg/mL. For gag-specific stimulation, 10 μL of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20 μL of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO<sub>2</sub>, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20 μL per tube anti-hCD3-APC, clone FN-18 (Biosource); 20 μL anti-hCD8-PerCP, clone SK1 (Becton Dickinson);

and 20  $\mu$ L anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750  $\mu$ L 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in PBS/2%FBS and 0.1  $\mu$ g of FITC-anti-hIFN- $\gamma$ , clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence cut-off for cytokine-positive events was used for both CD4<sup>+</sup> and CD8<sup>+</sup> populations, and for both mock and gag-peptide reaction tubes of a sample.

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#### D. Anti-p24 ELISA

A modified competitive anti-p24 assay was developed using reagents from the Coulter p24 Antigen Assay kit (Beckman Coulter, Fullerton, CA). Briefly, to a 250-μL serum sample, 20 μL of Lyse Buffer and 15 μL of p24 antigen (9.375 pg) from the Coulter kit were added. After mixing, 200 μL of each sample were added to wells coated with a mouse anti-p24 mAb from the Coulter kit and incubated for 1.5 hr at 37°C. The wells were then washed and 200 μL of Biotin Reagent (polyclonal anti-p24-biotin) from the Coulter kit was added to each well. After a 1 hr, 37°C incubation, detection was achieved using strepavidin-conjugated horseradish peroxidase and TMB substrate as described in the Coulter Kit. OD450nm values were recorded. A 7-point standard curve was generated using a serial 2-fold dilution of serum from an HIV-seropositive individual. The lower cut-off for the assay is arbitrarily set at 10 milli Merck units/mL (mMU/mL) defined by a dilution of the seropositive human serum. This cutoff falls at approximately 65% of the maximum bound control signal which corresponds to that obtained with the diluent control only and with no positive analyte.

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#### E. Results

PBMCs collected at regular 4-wk intervals were analyzed in an ELISPOT assay (Figure 17). Both Ad24ΔE1gagΔOrf6Ad5Orf6 and Ad24ΔE1gagΔE4Ad5Orf6 were able to induce significant levels of gag-specific T cells in non-human primates. At 10^11 vp dose level, the Ad24-induced responses were within 2-3-fold of those of MRKAd5-HIV1 gag. Both Ad24 vectors were also able to induce detectable levels of gag-specific T cells at 10^10 vp but were lower than those observed using MRKad5gag at the same dose.

PBMCs collected at wk 12 from the vaccinees were analyzed for intracellular IFN-γ staining after the priming immunizations. The assay results provided information on the relative amounts of CD4<sup>+</sup> and CD8<sup>+</sup> gag-specific T cells in the peripheral blood (Figure 18). The

results indicated that the prime-boost immunization approach was able to elicit in rhesus macaques both HIV-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cells.

#### F. Humoral Immune Responses

The Ad24-based vaccine vector was able to generate detectable levels of circulating anti-gag antibodies at the reasonably high dose level (Figure 19). No detectable titers were observed at equal to or lower than 10^10 vp, suggesting the existence of a dose-dependent response.

#### EXAMPLE 14

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In Vivo Transgene Expression

#### A. Immunization

Cohorts of 5 C3H/HeN mice were given single intramuscular injections of one of the following vectors: (1) 10^10 vp Ad24ΔE1SEAPΔE4Ad5Orf6; (2) 10^10 vp Ad24ΔE1SEAPΔOrf6Ad5Orf6; (3) 10^10 vp MRKAd5SEAP; and (4) 10^9 vp MRKAd5SEAP. Female mice were between 4-10 weeks old. The total dose of each vaccine was suspended in 0.1 mL of buffer. The vectors were given to both quadriceps of each of the animals with a volume of 50 uL per quad and using 0.3-mL 28G1/2 insulin syringes (Becton-Dickinson, Franklin Lakes, NJ). For the primates, the total dose of each vaccine was suspended in 1 mL of buffer. The monkeys were anesthetized (ketamine/xylazine mixture) and the vaccines were delivered i.m. in 0.5-mL aliquots into two muscle sites using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Serum samples were collected at defined intervals and stored frozen until the assay date. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council.

#### B. SEAP Assay

Serum samples were analyzed for circulating SEAP levels using TROPIX phospha-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5 uL aliquots of each serum were mixed with 45 uL of kit-supplied dilution buffer in a 96-well white DYNEX plate. Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the wells for 30 minutes at 65 °C.

Enzymatic SEAP activities in the samples were determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorder using DYNEX luminometer. RLU readings are converted to ng/mL SEAP using a log-log regression analyses.

#### 5 C. Rodent Results

Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 20. Results indicate that (1) both Ad24 constructs are all capable of expressing the SEAP transgene in vivo to comparable levels; and that (2) the level of expression achieved using the Ad24 vectors are comparable to that of Ad5 at 10-fold lower dose. The levels of SEAP in the serum dropped dramatically after day 2 and were at background levels by day 12.

#### D. Primate Results

Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the following vectors: (1) 10^11 vp MRKAd5-SEAP; (2) 10^9 vp MRKAd5-SEAP; (3) 10^11 vp Ad24ΔE1SEAPΔOrf6Ad5Orf6; or (4) 10^11 vp Ad24ΔE1SEAPΔE4Ad5Orf6. Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 21.

Results indicate that the peak levels of SEAP product produced by adenovirus serotype 24 were lower than but were within 3-fold of that of MRKAd5 at the same high dose level of 10^11 vp (Figure 21). The levels observed with adenovirus serotype 24 are generally 50-fold higher than those observed using 10^9 vp of MRKAd5. The levels of SEAP in the serum dropped dramatically after day 10 and were close to background as early as day 15. These observations strongly indicate that adenovirus serotype 24 is very efficient in expressing a transgene following intramuscular administration in a primate.

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#### **EXAMPLE 15**

# Construction of pMRKAd24ΔE1ΔE4Ad5Orf6

To construct pMRKAd24 $\Delta$ E1 $\Delta$ E4Ad5Orf6 (An Ad24 pre-Ad plasmid, composed entirely of Ad24 sequence and containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), an Ad24 ITR cassette was constructed containing sequences from the right (bp 31978 to 32264 and bp 34713 to 35164) and left (bp 4 to 450 and bp 3364 to 3799) end of the Ad24 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd24-4. Next the Ad5 Orf6 open reading frame (Ad5 bp 31192 to bp 34078) was generated by PCR and cloned between Ad24 bp 32264 and 34713 generating

pNEBAd24E-Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad24 bp 451 to 3363 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad24 bp 32265 to 34712 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct Ad5 Orf6 expression is driven by the Ad24 E4 promoter. The Ad24 sequences (bp 31978 to 32264 and bp 3464 to 3799) in the ITR cassette provide regions of homology with the purified Ad24 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria (Figure 22). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion will release the recombinant Ad24 genome from plasmid sequences. Potential clones will be screened by restriction analysis and one clone was selected as pMRKAd24ΔE1ΔE4Ad5Orf6. Pre-Adenovirus plasmid pMRKAd24ΔE1ΔE4Ad5Orf6 should contain Ad24 sequences from bp 4 to 450; bp 3364 to bp 32264 and bp 34713 to bp 35164 with Ad5Orf6 cloned between bp 32264 and bp 34713. The bp numbering in the above description refers to the wt sequence for both Ad24 and Ad5.

#### **EXAMPLE 16**

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## Insertion of HIV-1 gag and SEAP transgenes into pAd24ΔE1ΔE4Ad5Orf6

In order to introduce a gag or SEAP expression cassettes into the E1 region of pMRKAd24ΔE1ΔE4Ad5Orf6, bacterial recombination will be used. An HIV-1 gag expression cassette will consist of the following: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence, in the E1 deletion of an Ad24 shuttle plasmid, pNEBAd24-2 (a precursor to the Ad24 ITR cassette described above), generating pNEBAd24CMVgagBGHpA. PNEBAd24-2 contains Ad24 sequences from the left end of the genome (bp 4 to 450 and bp 3364 to 3799) that define the E1 deletion. The gag expression cassette will be obtained from a previously constructed plasmid and cloned into the E1 deletion between bp 450 and 3364 in the E1 parallel orientation. The shuttle vector containing the gag transgene will be digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad24 bp 4 to 450 and bp 3364 to 3799 and the fragment will be purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pMRKAd24ΔE1ΔE4Ad5Orf6 which was linearized in the E1 region by digestion with SwaI, should result in the generation of Ad24 gag-

containing pre-Adenovirus plasmids pMRKAd24ΔE1gagΔE4Ad5Orf6 by homologous recombination. Potential clones will be screened by restriction analysis.

A similar strategy will be used to generate Ad24 pre-Ad plasmids containing a SEAP expression cassette. In this case, a SEAP expression cassette will consist of: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence cloned into the E1 deletion of an Ad24 shuttle plasmid, pNEBAd24-2, generating pNEBAd24CMVSEAPBGHpA. The transgene will then be recombined into pMRKAd24 $\Delta$ E1 $\Delta$ E4Ad5Orf6 as described above for the gag transgene.

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# EXAMPLE 17 In Vivo Immunogenicity

#### A. Immunization

Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the *Guide for Care and Use of Laboratory Animals*, Institute of Laboratory Animal Resources, National Research Council.

#### 25 B. T Cell Responses

Ad24 Vaccine Vector as a Heterologous Booster: Cohort of 4 rhesus macaques was immunized initially with 3 doses (wk 0, 4, 26) of either 10<sup>7</sup> or 10<sup>9</sup> vp of MRKAd5-gag (see, PCT/US01/28861, published March 21, 2002) or MRKAd6-gag. At wk 56, the animals received a booster vaccine of 10<sup>11</sup> vp Ad24ΔE1gagΔOrf6Ad5Orf6. A separate cohort of naīve animals received a single dose of the booster vaccine. The results of the IFN-γ ELISPOT analyses of PBMC collected during the course of the studies are shown in Figure 23. It is apparent that the Ad24 HIV vectors can be utilized to amplify the existing pools of HIV-specific T cells. The increases in the levels of gag-specific T cells from the pre-boost levels to those measured at 4 wks post boost were consistently larger than the levels induced by the same booster vaccine in naïve animals. PBMCs from the vaccinees of the heterologous MRKAd5/MRKAd6-Ad24 boost

regimen were analyzed for intracellular IFN-γ staining after the priming immunizations (wk 60). The assay results provided information on the relative amounts of CD4<sup>+</sup> and CD8<sup>+</sup> gag-specific T cells in the peripheral blood (Figure 24). The results indicated that heterologous prime-boost immunization approach was able to elicit in rhesus macaques both HIV-specific CD4+ and CD8+ T cells.

Ad24 Vaccine Vector as a Heterologous Primer: In a separate study, a cohort of 3 rhesus macaques was immunized initially with 2 doses (wk 0, 4) of 10<sup>11</sup> vp Ad24ΔE1gagΔOrf6Ad5Orf6 and boosted at wk 24 with 10<sup>7</sup> vp of MRKAd5-gag. The low dose of MRKAd5-gag is selected to mimic the effect of pre-existing neutralizing immunity to the vector in a subject. A separate cohort of naïve animals was given a single dose of 10<sup>7</sup> vp MRKAd5-gag. The results of the IFN-γ ELISPOT analyses of PBMC collected during the course of the studies are shown in Figure 25.

The Ad24-based vaccine was able to prime effectively for HIV-specific T cell responses in macaques. Boosting with a low dose MRKAd5-gag resulted in a significant increase in the levels of gag-specific T cells. The increases in 2 out of 3 animals exceed the levels typically observed after treatment of naïve animals with the same low dose of MRKAd5-gag.

#### EXAMPLE 18

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#### 20 Construction of pAd34ΔE1ΔE4Ad5Orf6

To generate an E1- Ad34 based vector that can propagate in existing group C/Ad5 E1 complementing cell lines (293, PER.C6), Ad5 Orf6 was inserted in place of the native E4 region. Since at the time, the complete sequence of Ad34 (see Figures 28A-1 to 28A-9; subject of copending application serial no. 60/458,825, filed March 28, 2003) was unknown, advantage was taken of the sequence homology between Ad34 and Ad35 in order to construct the Ad34 pre-Adenovirus plasmid. Cotransformation of BJ 5183 bacteria with purified wild-type Ad34 viral DNA and the appropriately constructed Ad35 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The construction of the pre-Ad plasmid based on Ad34, is outlined below:

To construct pAd34ΔE1ΔE4Ad5Orf6 (An Ad34 pre-Ad plasmid containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), we utilized an Ad35 ITR cassette. We anticipated that sequence homology between Ad34 and Ad35 would allow homologous recombination to occur. The Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31913 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome (see Figures 2A-1 to 2A-10) separated by plasmid sequences containing a

bacterial origin of replication and an ampicillin resistance gene. The four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 31913 and 34419 generating pNEBAd35-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31914 to 34418 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31913 and bp 3403 to 3886) in the ITR cassette provided regions of homology with the purified Ad34 viral DNA in which bacterial recombination could occur following cotransformation into BJ 5183 bacteria (Figure 26). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion would release the recombinant Ad34 genome from the plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd34ΔE1ΔE4Ad5Orf6.

#### **EXAMPLE 19**

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### 20 Rescue of pAd34ΔE1ΔE4Ad5Orf6 into Virus

In order to determine if pre-adenovirus plasmid pAd34\Delta E1\Delta E4Ad5Orf6, could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique (Cell Phect Transfection Kit, Amersham Pharmacia Biotech Inc). *Pme*I digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring was observed following transfection. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation.

Approximately 1 ml of the cell lysate was used to infect a T-225 flask of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following complete CPE, the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment

followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with *HindIII* and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with *Pme1/HindIII* prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

#### EXAMPLE 20

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### Insertion of an Expression Cassette into pAd34ΔE1ΔE4Ad5Orf6

In order to introduce a gag or SEAP expression cassette (see Figures 6 and 7, respectively) into the E1 region of pAd34ΔE1ΔE4Ad5Orf6, bacterial recombination was again used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence, was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2 (a precursor to the Ad35 ITR cassettes described above), generating pNEBAd35CMVgagBGHpA. pNEBAd35-2 contains Ad35 sequences from the left end of the genome (bp 4 to 456 and bp 3403 to 3886) with a unique SwaI site between bp 456 and 3403 at the position of the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pNEBAd35-2. This cloning step resulted in the gag expression cassette being inserted into the E1 deletion between bp 456 and 3403 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad35 bp 4 to 456 and bp 3403 to 3886 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pAd34AE1AE4Ad5Orf6, linearized in the E1 region by digestion with Swa I, resulted in the generation of the Ad34 gag-containing pre-Adenovirus plasmid pAd34AE1gagAE4Ad5Orf6 by homologous recombination. Potential clones were screened by restriction analysis.

A similar strategy was used to generate Ad34 pre-Ad plasmids containing a SEAP expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence was

cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2, generating pNEBAd35CMVSEAPBGHpA. The SEAP expression cassette was obtained from a previously constructed shuttle plasmid by *Eco*RI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the *Swa*I site in pNEBAd35-2. The transgene was then recombined into the pAd34ΔE1ΔE4Ad5Orf6, generating pAd34ΔE1SEAPΔE4Ad5Orf6 as described above for the gag transgene.

All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

#### 10 EXAMPLE 21

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# Construction of pMRKAd34\Delta E1\Delta E4Ad5Orf6

To construct an Ad34 pre-Ad plasmid that was composed entirely of Ad34 sequences, an Ad34 ITR cassette was generated. The Ad34 ITR cassette was constructed containing sequences from the right (bp 31584 to 31895 and bp 34409 to 34772) and left (bp 4 to 456 and bp 3402 to 3885) end of the Ad34 genome (see Figures 28A-1 to 28A-9) separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd34-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad34 bp 31895 and 34409 generating pNEBAd34-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad34 bp 457 to 3401 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad34 bp 31896 to 34408 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct Ad5Orf6 expression is driven by the Ad34 E4 promoter. The Ad34 sequences (bp 31584 to 31895 and bp 3402 to 3885) in the ITR cassette provided regions of homology with the purified Ad34 viral DNA in which bacterial recombination could occur following cotransformation into BJ 5183 bacteria (Figure 27). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion would release the recombinant Ad34 genome from the plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pMRKAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

# EXAMPLE 22 In Vivo Studies

#### A. Immunization

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Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the two vectors: (1) 10^11 vp MRKAd5-SEAP (in MRKAd vector backbone disclosed in PCT/US01/28861, published March 21, 2002); and (2) 10^11 vp Ad34ΔE1SEAPΔE4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council.

#### B. SEAP Assay

Serum samples were analyzed for circulating human secreted alkaline phosphatase (SEAP) levels using TROPIX phospha-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5 µL aliquots of each serum were mixed with 45 µL of kit-supplied dilution buffer in a 96-well white DYNEX plate. Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the well for 30 minutes at 65 °C. Enzymatic SEAP activities in the samples were determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorded using DYNEX luminometer. RLU readings were converted to ng/mL SEAP using a log-log regression analyses.

#### C. ELISPOT Assay

The IFN- $\gamma$  ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen *et al.*, 2001 *J. Virol.* 75(2):738-749), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50  $\mu$ L of 2-4 x 10<sup>5</sup> peripheral blood mononuclear cells (PBMCs) were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower

size cut-off set at 80 femtoliters ("fL"). Either 50  $\mu$ L of media or the gag peptide pool at 8  $\mu$ g/mL concentration per peptide was added to the PBMC. The samples were incubated at 37°C, 5% CO<sub>2</sub> for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform (Silver Spring, MD); the counts were normalized to  $10^6$  cell input.

# D. Intracellular Cytokine Staining (ICS)

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To 1 ml of 2 x 106 PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added to a final concentration of 1  $\mu$ g/mL. For gag-specific stimulation, 10  $\mu$ L of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20  $\mu L$  of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO<sub>2</sub>, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20 μL per tube antihCD3-APC, clone FN-18 (Biosource); 20 μL anti-hCD8-PerCP, clone SK1 (Becton Dickinson); and 20 µL anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750  $\mu$ L 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in PBS/2%FBS and 0.1 μg of FITC-anti-hIFN-γ, clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence cut-off for cytokine-positive events was used for both CD4<sup>+</sup> and CD8<sup>+</sup> populations, and for both mock and gag-peptide reaction tubes of a sample.

#### E. Results

Expression: Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 29. Results indicate that the peak levels of SEAP protein produced by the alternative adenovirus serotype were lower than but were within 3-fold of that of MRKAd5 at the same high dose level of 10^11 vp (Figure 29). The levels of SEAP in the serum dropped dramatically after day 10 and were close to background as early as day 15. These observations strongly indicate that the Ad34-based vector is efficient in expressing a transgene following intramuscular administration in a primate.

Immunogenicity: Vaccine-induced T cell responses against HIV-1 gag were quantified using IFN-gamma ELISPOT assay against a pool of 20-aa peptides that encompassed the entire protein sequence. The results are shown in Figure 30; they are expressed as the number of spot-forming cells (SFC) per million peripheral blood mononuclear cells (PBMCs) that responded to the peptide pool or the mock (no peptide) control.

Immunization with gag-expressing Ad34 vector induced detectable levels of circulating gag-specific T cells immediately after a single dose of the vector. The responses improved following a second dose given at wk 4. Overall, the responses to the Ad34-based vector were slightly lower than those induced by the same dose of MRKAd5-gag. The results strongly indicate the Ad34-based vector can prime effectively for HIV-specific T cell responses.

IFN-γ ICS analyses of the PBMC from the Ad34-immunized animals revealed that the vector can induce detectable levels of both CD4<sup>+</sup> and CD8<sup>+</sup> HIV-specific T cells (Figure 31).

#### EXAMPLE 23

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#### Heterologous Immunization

Cohorts of 3 monkeys were immunized (at wks 0, 4) with 10^11 vp Ad34\Delta E1gag\Delta E4Ad5Orf6 followed by a booster at week 24 with 10^10 vp Ad35\Delta E1gag\Delta E4Ad5Orf6. Vaccine-induced T cell responses against HIV-1 gag were quantified using IFN-gamma ELISPOT assay against a pool of 20-aa peptides that encompassed the entire protein sequence. The results are shown in Figure 32; they are expressed as the number of spot-forming cells (SFC) per million peripheral blood mononuclear cells (PBMCs) that responded to the peptide pool or the mock (no peptide) control.

Immunization with gag-expressing Ad34 vector induced detectable levels of circulating gag-specific T cells that decreased to between 94-139 SFC/10^6 PBMC at the time of the boost. Heterologous immunization with an Ad35-based HIV vector resulted in as much as a 3-fold increase in T cell responses.

IFN-γ ICS analyses of the PBMCs from the Ad34 primed/Ad35 boosted animals at week 28 revealed that the vector can induce detectable levels of both CD4<sup>+</sup> and CD8<sup>+</sup> HIV-specific T cells (Figure 33).

#### WHAT IS CLAIMED IS:

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1. A means for propagating replication-defective adenovirus in an adenoviral E1-complementing cell line expressing E1 gene product(s) which are non-native to the adenovirus, which comprises:

- (a) inserting all or a portion of a heterologous adenoviral E4 region comprising nucleic acid sequence encoding open reading frame 6 (ORF6) into a replication-defective adenovirus; wherein the E4 region or portion thereof inserted into the adenovirus is native to a virus of the same adenovirus serotype as the E1 gene product(s) expressed by the complementing cell line;
- (b) introducing the replication-defective adenovirus into the adenoviral E1-complementing cell line;
- (c) allowing the replication-defective adenovirus to propagate in the adenoviral E1-complementing cell line; and
  - (d) rescuing the propagated adenovirus.
- 2. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof comprises the complete adenoviral E4-encoding region.
- 3. A means in accordance with claim 2 wherein the heterologous adenoviral E4 region or portion thereof comprises the complete adenoviral E4-encoding region and native E4 promoter.
  - 4. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof is inserted into the replication-defective virus in place of nucleic acid sequence encoding open reading frame 6 (ORF6).

5. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof is inserted into the replication-defective virus in place of nucleic acid sequence encoding the complete adenoviral E4-encoding region.

- 6. A means in accordance with claim 1 wherein the heterologous adenoviral

  5 E4 region or portion thereof is derived from a subgroup C adenovirus.
  - 7. A means in accordance with claim 1 wherein the subgroup C adenovirus is adenovirus of serotype 5.
  - 8. A means in accordance with claim 7 wherein the replication-defective adenovirus is an adenovirus of subgroup B.
  - 9. A means in accordance with claim 7 wherein the replication-defective adenovirus is an adenovirus of serotype 35.

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- 10. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof is operatively linked to a heterologous promoter.
- 11. A means in accordance with claim 1 wherein the adenoviral E1-complementing cell line is a PER.C6® cell line.
- 12. A replication-defective adenovirus comprising all or a portion of a heterologous E4 region comprising a heterologous adenoviral open reading frame 6 (ORF6).
- 13. A replication-defective adenovirus in accordance with claim 12 wherein the adenovirus comprises a heterologous gene of interest.
- 14. A replication-defective adenovirus in accordance with claim 13 wherein the heterologous gene of interest is a gene encoding an HIV-1 antigen.
- 15. A replication-defective adenovirus in accordance with claim 14 wherein the HIV-1 antigen is selected from the group consisting of HIV-1 gag, pol, nef and env.

16. A replication-defective adenovirus comprising all or a portion of a heterologous E4 region comprising a heterologous adenoviral open reading frame 6 (ORF6) and a gene encoding HIV-1 gag.

- 17. A replication-defective adenovirus comprising all or a portion of a

  beterologous E4 region comprising a heterologous adenoviral open reading frame 6 (ORF6) in

  place of a native E4 region or portion thereof comprising ORF6.
  - 18. A replication-defective adenovirus comprising all or a portion of a heterologous E4 region comprising a complete heterologous E4 region in place of a complete native E4 region.

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- 19. A replication-defective adenovirus comprising a heterologous E4 region or portion thereof comprising a complete heterologous E4 region including E4 promoter in place of a complete native E4 region.
  - 20. Adenovirus propagated in accordance with the means of claim 1.
- 21. A means in accordance with claim 1 wherein the replication-defective adenovirus comprises a heterologous gene of interest.
  - 22. A means in accordance with claim 21 wherein the heterologous gene of interest is a gene encoding an HIV-1 antigen.
  - 23. A means in accordance with claim 22 wherein the HIV-1 antigen is selected from the group consisting of: HIV-1 gag, pol, nef and env.
  - 24. A replication-defective adenovirus of serotype 35 comprising all or a portion of an adenovirus serotype 5 E4 region comprising open reading frame 6 (ORF6) and a heterologous gene of interest.
  - 25. A replication-defective adenovirus in accordance with claim 24 wherein the heterologous gene of interest is a gene encoding an HIV-1 antigen.

26. A replication-defective adenovirus in accordance with claim 25 wherein the HIV-1 antigen is selected from the group consisting of: HIV-1 gag, pol, nef and env.

27. A replication-defective adenovirus of serotype 35 comprising all or a portion of an adenovirus serotype 5 E4 region comprising open reading frame 6 (ORF6) and a gene encoding HIV-1 gag.

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- 28. A recombinant adenoviral vector of serotype 24 which comprises an E4 gene or a segment of an E4 gene comprising open reading frame 6 ("ORF6") of an alternative serotype.
- 29. A population of cells comprising the recombinant adenoviral vector of claim 28.
  - 30. A method for producing recombinant, replication-defective adenovirus particles comprising:
  - (a) introducing a recombinant adenoviral vector of claim 28 into a population of cells expressing adenovirus E1; and
    - (b) harvesting the resultant recombinant, replication-defective adenovirus.
  - 31. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 30.
  - 32. A composition comprising purified recombinant adenovirus particles in accordance with claim 31.
- 33. A composition in accordance with claim 32 which comprises a physiologically acceptable carrier.
- 34. A recombinant adenoviral vector in accordance with claim 28 which is at least partially deleted in E1 and devoid of E1 activity and comprises a heterologous nucleic acid.

35. A composition comprising purified recombinant adenoviral particles in accordance with claim 31 which are at least partially deleted in E1 and devoid of E1 activity and comprise a heterologous nucleic acid.

36. A method for effecting the delivery and expression of heterologous nucleic acid comprising administering the composition of claim 35 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.

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- 37. A method in accordance with claim 36 wherein the composition is preceded or followed by administration of heterologous nucleic acid with an adenovirus of a different serotype.
- 38. A composition in accordance with claim 35 wherein the heterologous nucleic acid encodes an HIV antigen.
- 39. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 38.
- 40. A composition in accordance with claim 39 wherein the HIV antigen is

  HIV-1 gag or immunologically relevant modification thereof.
  - 41. A composition in accordance with claim 39 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
  - 42. A composition in accordance with claim 39 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.
  - 43. A recombinant adenoviral vector of serotype 24 which is at least partially deleted in E1 and devoid of E1 activity; wherein said vector comprises an E4 gene or a segment of an E4 gene from adenovirus serotype 5 comprising open reading frame 6 ("ORF6"), and a heterologous nucleic acid.

44. A population of cells comprising the recombinant adenoviral vector of claim 43.

- 45. A method for producing recombinant, replication-defective adenovirus particles comprising:
- (a) introducing a recombinant adenoviral vector of claim 43 into a population of cells expressing adenovirus serotype 5 E1; and

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- (b) harvesting the resultant recombinant, replication-defective adenovirus.
- 46. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 45.
- 47. A composition comprising purified recombinant adenovirus particles in accordance with claim 46.
- 48. A composition in accordance with claim 47 which comprises a physiologically acceptable carrier.
- 49. A method for effecting the delivery and expression of the heterologous nucleic acid comprising administering the composition of claim 48 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.
- 50. A method in accordance with claim 49 above wherein the composition is preceded or followed by administration of the heterologous nucleic acid with an adenovirus of a different serotype.
- 51. A composition in accordance with claim 48 wherein the heterologous nucleic acid encodes an HIV antigen.
- 52. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 51.

53. A composition in accordance with claim 51 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.

- 54. A composition in accordance with claim 51 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
- 55. A composition in accordance with claim 51 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.

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- A recombinant adenoviral vector of serotype 34 which comprises an E4 gene or a segment of an E4 gene comprising open reading frame 6 ("ORF6") of an alternative serotype.
- 57. A population of cells comprising the recombinant adenoviral vector of claim 56.
- 58. A method for producing recombinant, replication-defective adenovirus particles comprising:
- (a) introducing a recombinant adenoviral vector of claim 56 into a population of cells expressing adenovirus E1; and
  - (b) harvesting the resultant recombinant, replication-defective adenovirus.
  - 59. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 58.
- 60. A composition comprising purified recombinant adenovirus particles in accordance with claim 59.
  - 61. A composition in accordance with claim 60 which comprises a physiologically acceptable carrier.
  - 62. A recombinant adenoviral vector in accordance with claim 56 which is at least partially deleted in E1 and devoid of E1 activity and comprises a heterologous nucleic acid.

63. A composition comprising purified recombinant adenoviral particles in accordance with claim 59 which are at least partially deleted in E1 and devoid of E1 activity and comprise a heterologous nucleic acid.

64. A method for effecting the delivery and expression of heterologous nucleic acid comprising administering the composition of claim 63 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.

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- 65. A method in accordance with claim 64 wherein the composition is preceded or followed by administration of heterologous nucleic acid with an adenovirus of a different serotype.
- 66. A composition in accordance with claim 63 wherein the heterologous nucleic acid encodes an HIV antigen.
- 67. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 66.
- 68. A composition in accordance with claim 67 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.
- 69. A composition in accordance with claim 67 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
- 70. A composition in accordance with claim 67 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.
- 71. A recombinant adenoviral vector of serotype 34 which is at least partially deleted in E1 and devoid of E1 activity; wherein said vector comprises an E4 gene or a segment of an E4 gene from adenovirus serotype 5 comprising open reading frame 6 ("ORF6"), and a heterologous nucleic acid.

72. A population of cells comprising the recombinant adenoviral vector of claim 71.

- 73. A method for producing recombinant, replication-defective adenovirus particles comprising:
- (a) introducing a recombinant adenoviral vector of claim 71 into a population of cells expressing adenovirus serotype 5 E1; and

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- (b) harvesting the resultant recombinant, replication-defective adenovirus.
- 74. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 73.
- 75. A composition comprising purified recombinant adenovirus particles in accordance with claim 74.
- 76. A composition in accordance with claim 75 which comprises a physiologically acceptable carrier.
- 77. A method for effecting the delivery and expression of the heterologous nucleic acid comprising administering the composition of claim 76 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.
- 78. A method in accordance with claim 77 above wherein the composition is preceded or followed by administration of the heterologous nucleic acid with an adenovirus of a different serotype.
- 79. A composition in accordance with claim 76 wherein the heterologous nucleic acid encodes an HIV antigen.
  - 80. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 79.

81. A composition in accordance with claim 79 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.

- 82. A composition in accordance with claim 79 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
- 5 83. A composition in accordance with claim 79 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.

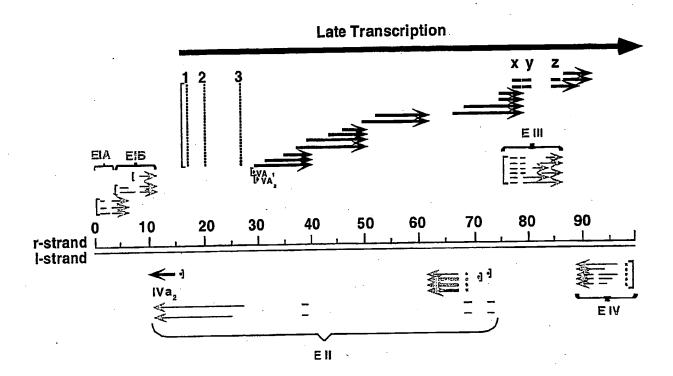


FIG. 1

1	catcatcaat	aatatacctt	atagatggaa	tggtgccaat	atgtaaatga	ggtgatttta
61	aaaagtgtgg	gccgtgtggt	gattggctgt	ggggttaacg	gttaaaaggg	geggegegge
121	cataggaaaa	tgacgtttta	tagagataga	attttttac	aagttgtcgc	gggaaatgtt
101	ogcoggaaaa	agacttettt	teteacquaa	ctacttagtt	ttcccacagt	atttaacagg
101	acycatadaa	aggettete		aaaattgctg	attttcccc	gaaaactgaa
241	aaatgaggta	gttttgaccg	gatgcaagtg	addattycty		ttettesea
301	tgaggaagtg	tttttctgaa	taatgtggta	tttatggcag	ggrggagrar	ttgttcaggg
361	ccaddtadac	tttgacccat	tacatagaga	tttcgattac	cgtgttttt	acctgaattt
121	ccacatacca	totcaaagto	ttctattttt	acgtaggtgt	cagctgatcg	ctagggtatt
401	tetegegeaceg	actttatata	sacacaccac	tcttgagtgc	садсдадаад	agttttctcc
481	tataccicag	ggtttgtgtt	aagaggccac	and an analytic	castttctac	ctcannaaat
541	tctgcgccgg	cagtttaata	ataaaaaaa	gagagatttg	-b	testeggaaa
601	aatctctgct	gagactggaa	atgaaatatt	ggagcttgtg	gigeacgeec	tyatyyyaya
661	cgatccggag	ccacctgtgc	agctttttga	gcctcctacg	cttcaggaac	tgtatgattt
721	agaggtagag	ggatcggagg	attctaatga	ggaagctgtg	aatggctttt	ttaccgattc
781	tatoctttta	actactaata	aaggattaga	attagatccg	cctttggaca	ctttcaatac
041	bacacacaca	attataaaa	acaatacaaa	tgtaagaaaa	ttacctgatt	tgagttccgt
841	tecaggggtg	accycygaaa	geggeacagg	atttestes	antratrarr	angaccatga
901	ggactgtgat	ttgcactgct	atgaagacgg	gtttcctccg	agtgatgagg	aggaccacga
961	aaaggagcag	tccatgcaga	ctgcagcggg	tgagggagtg	aaggergeea	atgutggttt
1021	tcagttggat	tacccaaaac	ttctggacat	ggctgtaagt	cttgtgaatt	tcacaggaaa
1081	aatactggag	taaaggaact	attatattca	cttttgttat	atgaaaaccc	actgccactt
11/1	tatttacact	aaagtgtgtt	taaottaaaa	tttaaaggaa	tatgctgttt	ttcacatgta
1201	***	and the tata	cttcttatta	taagtcctgt	atctgatgct	gatgaatcac
1201	caccyaytyt	bastcatages	tanactacta	atattcaagc	acctattcct	atagacatac
1261	cateteetga	Etetaetace	teaccidety	acattcaage	accegacee	goggacgogo
1321	gcaagcccat	tcctgtgaag	cttaagcctg	ggaaacgtcc	agcagcggag	aaacccgagg
1381	acttgttaca	gggtggggac	ggacctttgg	acttgagtac	acggaaacgt	ccaagacaat
1441	aagtgttcca	tatccgtgtt	tacttaaggt	gacgtcaata	tttgtgtgag	agtgcaatgt
1501	aataaaaata	tattaactat	tcactggttt	ttattgcttt	ttgggcgggg	actcaggtat
1561	ataantanaa	acadacctat	ataattaact	cataggagct	ggctttcatc	catggaggtt
1201	acaagcagaa	ternarior	taggeougee	aggcaactgt	tagagagggg	ttcggacgga
1071	Egggccattt	Lggaagacct	caygaagacc	aggeaactge	ctagagagaga	tttaggegge
1681	gtctccggtt	tttggagatt	etggtteget	agtgaattag	ctagggtagt	
1741	aaacaggact	ataaacaaga	atttgaaaag	ttgttggtag	attgcccagg	actititigaa
1.8.0.1	getettaatt	tgggccatca	ggttcacttt	aaagaaaaag	ttttatcagt	tttagacttt
1861	Fraaccccag	gtagaactgc	tactactata	gcttttctta	cttttatatt	agataaatgg
1001	atecceases	ctcatttcad	caggggatac	gttttggatt	tcatagccac	agcattgtgg
1741	accecycaga		caggggacac	atcttaggtt	actooccagt	gcagcctttg
TART	agaacatgga	aggiregeaa	yatyayyata	attactagget	caattataas	adaddaacad
2041	ggtgtagcgg	gaatcctgag	gcatccaccy	gtcatgccag	cggttctgga	ggaggaacag
2101	caagaggaca	acccgagagc	cggcctggac	cctccagtgg	aggaggcgga	gragergaer
2161	tgtctcctga	actgcaacgg	gtgcttactg	gatctacgtc	cactggacgg	gataggggcg
2221	ttaagaggga	gagggcatcc	agtggtactg	atgctagatc	tgagttggct	ttaagtttaa
2281	taaatcacaa	acotcctoaa	accatttoot	ggcatgaggt	tcagaaagag	ggaagggatg
2201	23agttgtag	attacaggan	asstattcac	tggaacaggt	gaaaacatgt	tggttggagc
2341	aaytttttgt	theresets	addatteeae	attatgccaa	gatagetttg	aggeetgata
2401	cagaggatga	rrgggaggrg	yccarraaaa	accargecaa be	ttettagett	teteesste
2461	aacagtataa	gatcagtaga	cggattaata	tccggaatgc	tryttacata	cciggaaatg
2521	gggctgaggt	ggtaatagat	actcaagaca	agacagttat	tagatgetge	atgatggata
2581	tataacctaa	agtagtcggt	atggaagcag	tcacttttgt	aaatgttaag	tttaggggag
2641	atoottataa	togaatagtg	tttatggcca	ataccaaact	tatattgcat	ggttgtagct
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2701		atattacatt	acceceacta	gcagaaccaa	gagtcaattg	tctctgaaga
2/01	gillicialyc	gructggact	gccacagccg	bb-be-be-b	342444444	accepted
2821	aatgcatatt	ccaaagatgt	aacctgggca	ttctgaatga	aggegaagea	agggtccgtc
2881	actgcgcttc	tacagatact	ggatgtttta	ttttaattaa	gggaaatgee	agegraaage
2941	ataacatgat	ttgtggtgct	tccgatgaga	ggccttatca	aatgctcact	tgtgctggtg
3001	ggcattgtaa	tatoctooct	actgtgcata	ttgtttccca	tcaacgcaaa	aaatggcctg
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3001	ttatagatta	caacgcgccg	atrastrato	tgaaagtgtt	attagaacca	gatgcctttt
3121	Clarycella	Coagrataac	atgaattatg	tgaaagtgee	aatataaaa	atectracet
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3241	atgatgatac	gagatcgagg	gtgcgcgcat	gcgaatgcgg	aggcaagcat	gecaggttee
3301	ageeggtgtg	tgtagatgtg	accgaagato	tcagaccgga	tcatttggtt	attgcccgca
3361	ctggagcaga	attcagatcc	agtggagaag	aaactgacta	aggtgagtat	tgggaaaact
3/21	ttaaaataaa	attttcacat	ggacagatto	agtaaaaatt	tatttttct	gtcttgcagc
2404 J#61	+ - 2 - 2 - 2 - 2 - 2 - 2 - 2 - 2 - 2 -	account	cttttaacco	gggagtcttc	agcccttatc	tgacaggggg
249T	Lyacatgagt	gyadatyttt	ttactcaayyy	, <u> </u>	totactotco	-sssacc
3541	tctcccatcc	rgggcaggag	cccagaa	tgttatggga	the activity	argyadyacc
3601	cgttcaaccc	gccaattctt	caacgctgac	ctatgctact	ttaagttctt	caccettgga
3661	cgcagctgca	gccgctgccg	ccgcctctgt	: cgccgctaac	actgtgcttg	gaatgggtta
3721	ctatggaage	atcgtggcta	attccacttc	ctctaataac	ccttctacac	tgactcagga
	2354					

			_			
3781	caagttactt	atecttttag	cccagctgga	ggctttgacc	caacgtctgg	gtgaactttc
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2041	Leageaggeg	geegageege	gagcacaac	teresee	5009900099	ttaageetaa
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3961	gtgtttttat	ttcatttttc	acacacaata	taccctagac	caccgatctc	gatcattgag
4001	202000000	atttttaaa	gaatggtata	gaggtgggat	tgaatgttta	gatacatggg
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1111	attataaata	acceatteat	aacaanntco	cantacataa	tgttgcacaa	tatcttttag
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4321	attgeegeca	agalecegic	LLyggillal	griacyaayy	actactaaga	cggcgcaccc
4381	ggtacattta	ggaaatttat	cgtgcagctt	ggatggaaaa	gcgtggaaaa	atttggagac
4441	accettatat	cctccgagat	tttccatgca	ctcatccato	ataatagcaa	tagaaccata
7277	acceeggg			et et en en en	tantaattat	~=====================================
4501	ggcagcggcg	egggcaaaca	egeteegegg	gictgatata	tcatagttat	greectgage
4561	taaatcatca	taagccattt	taatgaattt	ggggcggagc	gtaccagatt	ggggtatgaa
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28981	tgacttctgc	tcgctcacac	ctcactgtaa	tetataggete	taactgcatt	acaaaaccat
		. <u></u>	ccaccacccc	ocacaattat	. LLLLLLLUGG	ugcug cg ccg
		+	2 ** * * * * * * * * * * * * * * * * *	CAACAALCAU	Lateatere	googogoos
		- +~~+~~~++	cttadatttd	acacccaac		
			FOOTBCCEAU	agallicti		
			tracameant	acccacauca	accidadact	gtataggage
		• <b>+</b> -+~~~++	++actttat	TACELUCAL	Luculatuta	gualaguerg
			· aacttctada	ctadalccli	. ulucuaally	CCCaccageg
		, maatacccca	. accaaaatat	Cacaacact	. Ullayaulua	Lucadacua
		<b>- antannasta</b>		Tattuctic	: Clacquigu	ccaaccccay
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3006	1 gctggaatg	c tcccaatgca	catgatcato	. cacaayaccc	. ayayyaacac	. coccecat
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31801	aataaanttt	aagtgtttt	atttaaaatc	acaaaattcg	agtagttatt	ttgcctccac
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34081 34141 34201 34261 34321 34381 34441 34501 34561 34681 34681	tttccatcac gattaaacaa catacaatcc gtataattat gcacaggaga gtccctctaa ggcacacaaa gccctaaact ccgaaactgc ttcctcttc	aaacaagcgt aagacaagcc cagcaccgaa agacatgtta gcttaatcgt ataaaaaata atacacatac ccacaagctc gacgtaatgg gtcaccaggg tcaccggtacg tttaaccgtt ttacatattg	acagggtctc agttcctcgc gcatcagtta aagtatagca taattatttc aaagcctcat taaagtcact gactaaagtg acaagtacag acaagtacag	cagetegace ggtgaccage aggagaaaaa aagecacece tetgetgetg cagecatgge ctccaacete tataaattee tttaacttaca ccaatcacca	acagccaaca tcgcggatac tttaggcaac ttaccagaga tcaccaatat cgccaaaccc gcaatccaa acgtcatttt cacggccac	ctgtcatcgt cttgatgaag tagcctttgg aaagtaaaag gtcgccccg aagtacagcg atatacacaa aacaccacac caagcgtcac cccacggccg acttttaaa
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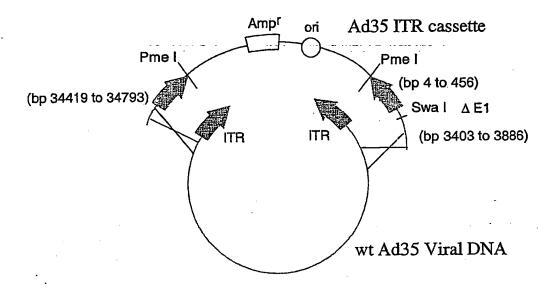


FIG. 3

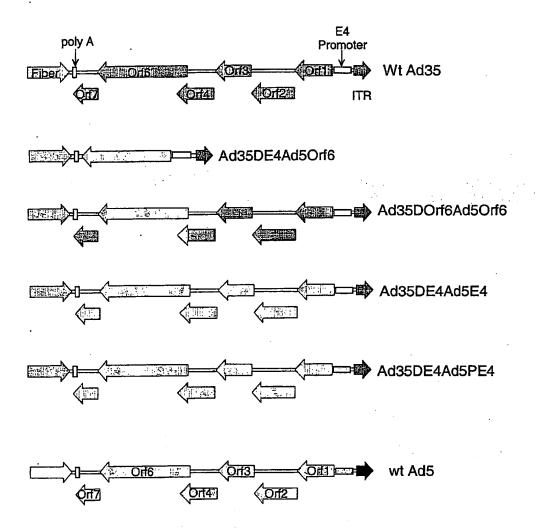
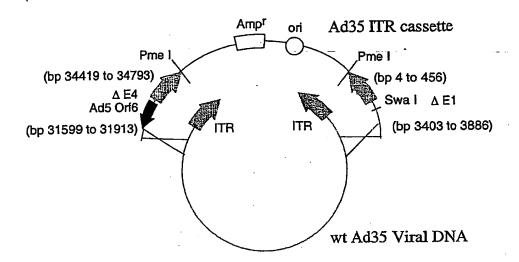


FIG. 4



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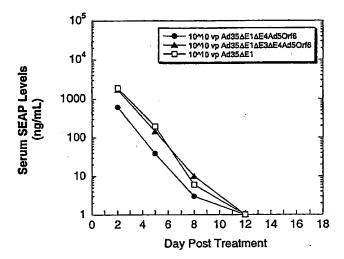
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121	ttagttcata	gcccatatat	ggagttccgc	gttacataac	ttacggtaaa	tggcccgcct
181	ggctgaccgc	ccaacgaccc	cegeceattg	acgtcaataa	tgacgtatgt	tcccatagta
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. 301	ttggcagtac	atcaagtgta	tcatatgcca	agtacgcccc	ctattgacgt	caatgacggt
361	aaatggcccg	cctggcatta	tgcccagtac	atgaccttat	gggactttcc	tacttggcag
421	tacatctacg	tattagtcat	cgctattacc	atggtgatgc	ggttttggca	gtacatcaat
481	gggcgtggat	agcggtttga	ctcacgggga	tttccaagtc	tccaccccat	tgacgtcaat
541	gggagtttgt	tttggcacca	aaatcaacgg	gactttccaa	aatgtcgtaa	caacteegee
601	ccattgacgc	aaatgggcgg	taggcgtgta	cggtgggagg	tctatataag	cagagetegt
661	ttagtgaacc	gtcagatcgc	ctggagacgc	catccacgct	gttttgacct	ccatagaaga
721	caccgggacc	gatccagcct	ccgcggccgg	gaacggtgca	ttggaacgcg	gattccccgt
781	gccaagagtg	_agatctaccA	TGGGTGCTAG	GGCTTCTGTG	CTGTCTGGTG	GTGAGCTGGA
841	CAAGTGGGAG	<b>AAGATCAGGC</b>	TGAGGCCTGG	TGGCAAGAAG	AAGTACAAGC	TAAAGCACAT
901	TGTGTGGGCC	TCCAGGGAGC	TGGAGAGGTT	TGCTGTGAAC	CCTGGCCTGC	TGGAGACCTC
961	TGAGGGGTGC	AGGCAGATCC	TGGGCCAGCT	CCAGCCCTCC	CTGCAAACAG	GCTCTGAGGA
1021	GCTGAGGTCC	CTGTACAACA	CAGTGGCTAC	CCTGTACTGT	GTGCACCAGA	AGATTGATGT
1081	GAAGGACACC	AAGGAGGCCC	TGGAGAAGAT	TGAGGAGGAG	CAGAACAAGT	CCAAGAAGAA
1141	GGCCCAGCAG	GCTGCTGCTG	GCACAGGCAA	CTCCAGCCAG	GTGTCCCAGA	ACTACCCCAT
1201	TGTGCAGAAC	CTCCAGGGCC	<b>AGATGGTGCA</b>	CCAGGCCATC	TCCCCCCGGA	CCCTGAATGC
1261	CTGGGTGAAG	GTGGTGGAGG	AGAAGGCCTT	CTCCCCTGAG	GTGATCCCCA	TGTTCTCTGC
1321	CCTGTCTGAG	GGTGCCACCC	CCCAGGACCT	GAACACCATG	CTGAACACAG	TGGGGGGCCA
1381	TCAGGCTGCC	ATGCAGATGC	TGAAGGAGAC	CATCAATGAG	GAGGCTGCTG	AGTGGGACAG
1441	GCTGCATCCT	GTGCACGCTG	GCCCCATTGC	CCCCGGCCAG	<b>ATGAGGGAGC</b>	CCAGGGGCTC
1501	TGACATTGCT	GGCACCACCT	CCACCCTCCA	GGAGCAGATT	GGCTGGATGA	CCAACAACCC
1561	CCCCATCCCT	GTGGGGGAAA	TCTACAAGAG	GTGGATCATC	CTGGGCCTGA	ACAAGATTGT
1621	GAGGATGTAC	TCCCCCACCT	CCATCCTGGA	CATCAGGCAG	GGCCCCAAGG	AGCCCTTCAG
1681	GGACTATGTG	GACAGGTTCT	ACAAGACCCT	GAGGGCTGAG	CAGGCCTCCC	AGGAGGTGAA
1741	GAACTGGATG	ACAGAGACCC	TGCTGGTGCA	GAATGCCAAC	CCTGACTGCA	AGACCATCCT
1801	GAAGGCCCTG	GGCCCTGCTG	CCACCCTGGA	GGAGATGATG	ACAGCCTGCC	AGGGGGTGGG
1861	GGGCCCTGGT	CACAAGGCCA	GGGTGCTGGC	TGAGGCCATG	TCCCAGGTGA	CCAACTCCGC
1921	CACCATCATG	ATGCAGAGGG	GCAACTTCAG	GAACCAGAGG	AAGACAGTGA	AGTGCTTCAA
. 1981	CTGTGGCAAG	GTGGGCCACA	TTGCCAAGAA	CTGTAGGGCC	CCCAGGAAGA	AGGGCTGCTG
2041	GAAGTGTGGC	AAGGAGGGCC	<b>ACCAGATGAA</b>	GGACTGCAAT	GAGAGGCAGG	CCAACTTCCT
2101	GGGCAAAATC	TGGCCCTCCC	ACAAGGGCAG	GCCTGGCAAC	TTCCTCCAGT	CCAGGCCTGA
2161	GCCCACAGCC	CCTCCCGAGG	AGTCCTTCAG	GTTTGGGGAG	GAGAAGACCA	CCCCCAGCCA
2221	GAAGCAGGAG	CCCATTGACA	AGGAGCTGTA	CCCCTGGCC	TCCCTGAGGT	CCCTGTTTGG
2281	CAACGACCCC	TCCTCCCAGT	<b>AA</b> aataaagc	ccgggcagat	ctgatctgct	gtgccttcta
2341	gttgccagcc	atctgttgtt	tgccctccc	cegtgeette	cttgaccctg	gaaggtgcca
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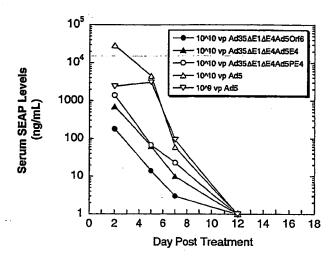
FIG. 6

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101	ggctgaccgc	CCSSCCSCCC	ccacccatta	acotcaataa	tgacgtatgt	teccatagta
241	acgccaatag	ggagtttcca	ttgacgtcaa	tagatagaat	atttaccota	aactgcccac
241	ttggcagtac	atgaagtgta	tcatatocca	agtacgcccc	ctattgacgt	caatgacggt
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301	-tacatctacg	tattagtgat	cactattacc	atgataataa	ggttttggca	gtacatcaat
421	gggcgtggat	accagicac	ctcaccacca	tttccaactc	tccaccccat	tgacgtcaat
481	gggagtttgt	tttgggtttga	asstcsacco	cactttccaa	aatgtcgtaa	caactcccc
541	ccattgacgc	222taggagga	taggggtgta	caataaaaa	tctatataag	cagagetegt
601	ttagtgaacc	adatgggcgg	ctggcgcgca	catccacact	attttaacct	ccatagaaga
551	caccgggacc	greagacege	ccggagacgc	gaacggtgca	ttagaacaca	gattccccgt
721	gccaagagtg	garceageer	cogoggeogg	CCTCCATCCT	CCTCCTCCTC	СТССТССТСС
781	GCCTGAGGCT	aga comence	CTCCCCATCA	TO COLOCATOCI	GCAGCAGAAC	CCGGACTTCT
841	GCCTGAGGCT	CCCACCCCCAC	CIGGGCAICA	CCCCCOTIGA	CCTCCACCCT	GCACAGACAG
901	CCGCCAAGAA	GGCAGCCGAG	WINCOUCCOCC	ATTCCCATGGG	CCTCTCTACC	CTCACACCTC
961	CCGCCAAGAA	CCTCATCATC	AACAACCACA	AIGGGAIGGG	TCACATACCC	CTGGCCATGG
1021	ACCGCTTCCC	AAAAGGGCAG	AMGAAGGACA COCOCCAACA	CAMACAAMOM	ACACAAACAT	CTCCCACACA
1081	GTGGAGCCAC	ATATGTGGCT	TA COMOMOCO	CATACAATGI	CAACAAACAI	ACCATTCCCT
1141	TGAGTGCAGC	AGCCACGGCC	AACCACTGCCA	ACACCACACC	CCCCAACCAC	CTCATCTCC
1201	TGAGTGCAGC	CGCCCGCTTT	AACCAGTGCA	ACACGACACG	CCCTACCACC	ACACCACTCC
1261	TGATGAATCG	GGCCAAGAAA	GCAGGGAAGT	LAGIGGGAGI	CCCCAACTCC	TACTCGAGIGC
1321	AGCACGCCTC CCGACGTGCC	GCCAGCCGGC	ACCTACGCCC	ACACGGIGAA	CCGCMACIGG	CACCOCACC
1381	CCGACGTGCC	TGCCTCCGCC	A MCCARGAGG	GGTGCCAGGA	CHICGCIACG	CCCATCCCAA
1441	CCAACATGGA	CATTGACGTG	ATCCTAGGTG	GAGGCCGAAA	GIACAIGIII	CACCCCAACA
1501	CCCCAGACCC	TGAGTACCCA	GATGACTACA	* COCHAGGIGG	GACCAGGCIG	A A C C C C A C T C
1561	ATCTGGTGCA	GGAATGGCTG	GCGAAGCGCC	AGGGTGCCCG	GIAIGIGIGG	MACCGCACIG
1621	AGCTCATGCA	GGCTTCCCTG	GACCCGTCTG	TGACCCATCT	CATGGGTCTC	AUCCACAUCA
1681	GAGACATGAA	ATACGAGATC	CACCGAGACT	CCACACTGGA	CCCCTCCCTG	ATGGAGATGA
1741	CAGAGGCTGC	CCTGCGCCTG	CTGAGCAGGA	ACCCCCCCCCG	CTTCTTCCTC	TICGIGGAGG
1801	GTGGTCGCAT	CGACCATGGT	CATCATGAAA	GCAGGGCTTA	CUGGGCACTG	ACTGAGACGA
1861	TCATGTTCGA	CGACGCCATT	GAGAGGGCGG	GCCAGCTCAC	CAGCGAGGAG	GACACGCTGA
1921	GCCTCGTCAC	TGCCGACCAC	TCCCACGTCT	TCTCCTTCGG	AGGCTACCCC	CTGCGAGGGA
1981	GCTCCATCTT	CGGGCTGGCC	CCTGGCAAGG	CCCGGGACAG	GAAGGCCTAC	ACGGTCCTCC
2041	TATACGGAAA	CGGTCCAGGC	TATGTGCTCA	AGGACGGCGC	CCGGCCGGAT	GTTACCGAGA
2101	GCGAGAGCGG	GAGCCCCGAG	TATCGGCAGC	AGTCAGCAGT	GCCCCTGGAC	GAAGAGACCC
2161	ACGCAGGCGA	GGACGTGGCG	GTGTTCGCGC	GCGGCCCGCA	GGCGCACCTG	GTTCACGGCG
2221	TGCAGGAGCA	GACCTTCATA	GCGCACGTCA	TGGCCTTCGC	CGCCTGCCTG	GAGCCCTACA
2281	CCGCCTGCGA	CCTGGCGCCC	CCCGCCGGCA	CCACCGACGC	CGCGCACCCG	GGTTAAcccg
2341	tggtccccgc	gttgcttcct	ctgctggccg	ggacatcagg	tggcccccgc	tgaattggaa
2401	tcgatcagaa	ttgatctgat	ctgc <u>tgtgcc</u>	ttctagttgc	cagccatctg	ttgtttgccc
2461	ctcccccgtg	ccttccttga	ccctggaagg	tgccactccc	actgtccttt	cctaataaaa
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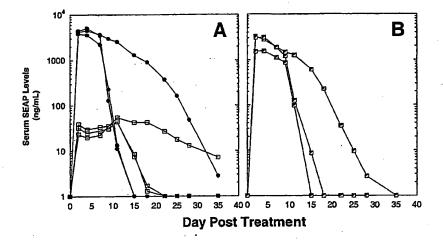
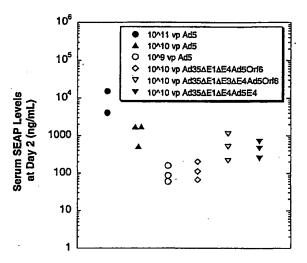


FIG. 10A-B



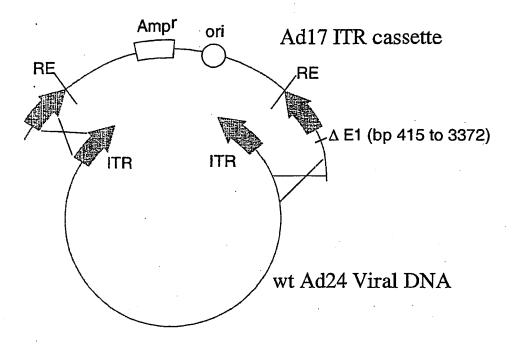
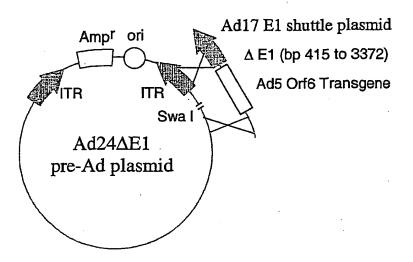


FIG. 12



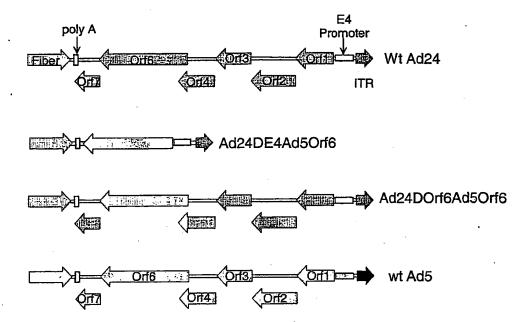
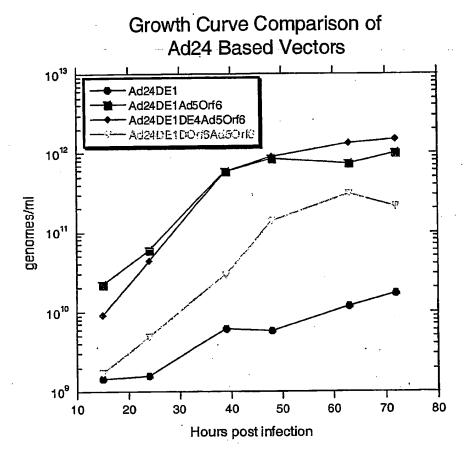


FIG. 14

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. 61	tttagggcgg	gaccagcact	gattggacga	gagaagatga	tgcaaatgac	gtcacgacgc
121	accordaacc	atcaccacaa	aggcataacc	tagcccggaa	gcaagtcgcg	gggctgatga
191	catataaaaa	accognacttt	agacccggaa	acggccgatt	ttcccacaac	cacqcccqqa
2/1	tatgaggtaa	ttctgggcgg	atgcaagtaa	aattaggtca	ttttaacaca	aaaactgaat
301	cacgaggtaa	aaantnaaaa	ataccootco	cgcccagggc	ggaatattta	ccaaaaacca
361	aggaagtga	accoattaco	tagaggtttc	gattgcggtg	tttttcaca	aatttcccc
401	tagagacticg	accepactace	ttatotcaca	gatcagctga	tccacagggt	atttaaacca
421	ctegigicaa	tanagecc	actetteact	gccagcgagt	agagatttct	ctgagctccg
401 E41	gregageeeg	ctaayayycc	tracacacct	gegeeteett	tetteaactq	tocctattoa
241 201	cittedagagt	ttattactaa	aggacacot	gagtacaata	ttggaggacg	aactgcatcc
C C 1	catggeegea	cactgorgg	ctacacttca	ggacctatat	gatttggagg	tagatgccca
721	tastasaasa	gagetgggae	aggetataaa	tttaatattt	ccagaatete	tgattcttca
701	ratesasta	ccgaacgaag	ctatacctac	accacttcat	acaccgácto	totcacccat
0/1	ggctgacata	gecagegaag	acrarctara	cctccgatgt	tatgaggaag	attttcctcc
041	accigaatig	gaagaggagg	acquircage	gagcatggct	ctaatctcaa	aatatootto
901	tatastata	gaggacgaac	ttatattaa	caatcctgag	atacccaaac	aaggctgtag
1001	rectagging	taccaccocc	ataagaccaa	agacacgaac	acetectaca	ctctgtgtta
1021	accongucay	actaccygy	ttatttacad	taagtggagt	gaatgtgaga	gagactgagt
11/1	catyaaaaay	taactcagct	atocttagac	agctgtgcta	agtgtggttt	atttttattt
1701	gerraacaca	tatacegggea	tractcatca	ccctcagaag	aadaccaccc	gtgtcccct
1201	eraggreegg	coccagagga	cctccaacta	cacagaccca	cccaatcaa	acceptage
1201	gagetgteag	gradatyce	assasttaaa	gacttgttac	atgacatgg	tagagatasa
1201	gagaggcgag	tagattass	addadccydy	aactaggctc	acctatactt	antcatotot
1201	cettiggate	ctagggggaa	actetatata	acgcatgcaa	agtataattt	atgactcatg
1441	aaataaaytt	gracaaraaa	agracacgeg	acctgggcac	tagaacacaa	accttcadd
1201	ggegeggeee	agicciatac	actatoctto	cagactttag	caagagacacag	caacttataa
1001 1001	ageteetgat	ggatgtgtgg	tecarettet	ggagacactg	atttaaaact	cctctatctc
1021	aggacagice	agacgggcgc	aaggattata	acgaggaatt	trasastrtt	tttactaatt
1001	gcctggtgta	catagitaay	ctasatctcc	gccaccagtc	ccttttccad	gaaagggtac
1001	teenggeet	tratttttca	accesace	gcactacagc	cagaattact	tttataattt
1001	ttetacayeee	casatagaaa	cacaacaccc	aactgagcag	agactacatt	ctagacttca
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1001	cagecatgea	cccgcggagg	ccaatcttc	ttcgtctaca	cacacaaaca	tccatattaa
1301	ggettataca	gecageagee	ataracaara	accegaggag	caacctaaac	cctccatcaa
2101	aggaagaaat	gaggcaggcc	acggacgaga	cctgtaccca	gagettagea	gaatactaac
2101	aayayyaycu	aggaccgaacc	aggeacecag	gagcgatggg	ageetagea	gggtgetgac
2701	acceatggee	aggggagtga	tastcass	gcgtccagag	cacattacct	ggatgatgat
2221	cyayetyacy	tatagaata	agatagaact	gatgcaggat	aaatataacc	tagaacagat
7201	acayarggag	tacttasec	caratrarra	ttgggaggag	accettaera	aatatoccaa
2341	aaaaacccac	caccagatt	cagacgagga	ggtgaccaag	accottoaata	tcagacatgo
2401	gatageeety	togggggacc	gcaagtacag	ggtcatcgat	accetagaca	aggccacctt
2401	cogctatatt	teggggaacg	tracarcorr	agtgatgaat	atraattrca	toattttcat
2521	caygrattaca	ttantagaa	agagette	tggggtgatg	ttcatccca	acadtcacat
220I	gaacatgaay	cccaacygay	tattagactt	caacaatatg	tacacagaca	tetagagagag
2041	yaccerycac	ggctgcagtt	actttata	ctgctggatg	accataatca	rearacces
2701	roccaagate	aggggatgta	agtetetet	tgagaaatgc	tacctooca	tctctacca
2/01	gagegagacg	cetgtgaage	agtgtgtgtt	cctggagacg	cactcottat	acctaatas
787T	gggcaatget	agagtgagac	attyctctt	cccggagacg	agecycece	gcctggtgaa
288I	gggcacagec	tetetgaage	acaacacggc	gaagggctgc	acggatgage	tracetecea
2941	catgetgaca	Lgegaetegg	gggtetgeca	tatcctgaag	adcatttaty	reatesaset
300I	ceeeeggaag	aagtggccag	togagagagagagagagagagagagagagagagagagaga	taacctactg	tttaggggg	ccaarctact
3061	gggcgccaga	aggggcacct	cccayccgta	ccagtgcaac	atatttaa.ca	tagatatata
3121	gctggagaac	gatgccttct	ccagggrgaa	cctgaacggc	atcouraged	aggacycccc
3181	ggtgtacaag	accetgagat	acgargagac	caagtccagg	gracegett	taaaaaaaa
3241	gggcagacac	accaggatge	aaccagtggc	cctggatgtg	accyaggage	ryayycccya
3301	ccacctggtg	atggcttgta	ccgggaccga	gttcagctcc	agugggagg	acacagatta
3361	gaggtaggtt	gagtattagt	gggcgcggct	aaggtgacta	LaaaggCggg	Lyttetacga
3421	gggtcttttt	gettttetge	agacatcatg	aacgggactg	geggggeett	cgaaggggg
3481	ctttttagcc	cttatttgac	aacccgcctg	ccgggatggg	ccggagttcg	cagaatgtg
3541	atgggatcga	cggtggacgg	gcgtccagtg	cttccagcaa	actectegae	catgacctac
3601	gcgaccgtgg	ggaactcgtc	acccaacaac	accgccgcag	ccgcggcagc	cycageegee

FIG. 16A-1

			**	-+	acaacaataa	cccctctctc
3661	atgacagcga	cgagactggc	ttegagetae	atgcccagca	gcagcagcag	ccccccgcg
3721	cccagttcca	tcatcgccga	ggagaaactg	ctggccctgc	tggccgagct	ggaagccctg
3781	addeddedade	taggegeet	gacccagcag	gtgtccgagc	tecgegaaca	gcagcagcag
3701		~attenataa	agagagattc	tgattcaaac	accasaccat	ctttattatt
3841	Cadadiadai	yarrcaaraa	acacagacce			atacastas
3901	tattttttcg	cgcgcggtag	geeetggtee	acctctcccg	accaccyaya	gracaara
3961	ttttttccag	gacccggtag	aggtgggatt	ggatgttgag	gtacatgggc	atgagcccgt
4021	cecaaaaata	gaggtaggag	cactocatoo	cctcgtgctc	taggategta	ttgtagatga
4001	teesetests	202200000	tagacataat	gctggatgat	gtccttgagg	aggagactga
408T	tecagicala	gcaggggcgc	cgggcgcgggc	beergaceac	attendates	aggagataca
4141	tggccacggg	gagccccttg	grgraggrgr	tggcgaagcg	grigagergg	yayyyacyca
4201	tgcgggggga	gatgatgtgg	agtttggcct	ggatcttgag	gttggcgatg	ttgccaccca
4261	gatecegect	ggggttcatg	ttgtgcagga	ccaccagaac	ggtgtagccc	gtgcacttgg
1221	greattata	atroaactto	gaagggaatg	cgtgaaagaa	tttggagacg	cccttatacc
4221	ggaactegte	tteesteese	testesstes	tgatggcgat	addeceataa	actacaactt
438T	cacccaggtt	ttecatgeac	leatecatga	Lyacyycyac	gggcccgcgg	acceptage
4441	tggcaaagac	gtttctgggg	tcagagacat	cgtaattatg	ereergggrg	ayattattat
4501	aagacatttt	aatgaatttg	gggcggaggg	tgccagattg	ggggacaatg	gttccctcgg
4561	uccccaaaac	gaagttcccc	tcacatattt	gcatctccca	ggctttcatc	tcggaggggg
4601	goodogggg	cacctacaaa	accetases	aaacggtttc	caaaacaaaa	gtgatgagct
4021	ggattatgtt		and a section	acttgccgca	cccaatcaa	ccatagatga
4681	gcgaggagag	caggiller	aacagcrygg	actigotyca	cccggccggg	teeeega
4741	ccccgatgac	gggttgcagg	tggtagttca	aggacatgca	getgeegteg	teceggagga
4801	ggggggcac	ctcgttgagc	atgtctctga	cttggaggtt	ttcccggacg	agctcgccga
1861	agaggggggg	cccacceaac	gagaggaggt	cttgcaggga	agcaaagttt	ttcaggggct
4001	tanagagge	accetacac	atcttggcga	gggtctgcga	gaggagttcg	aggcggtccc
4921	tgageeegte	ggccargggc	accedgega	gggtetgega	anthantant	ttaaaaaatt
4981	agagctcggt	gacgtgctct	acggcatete	gatccagcag	acticcicgt	cccgggggcc
5041	gggacgactg	cgactgtagg	gcacgagacg	atgggcgtcc	agcgctgcca	gcgtcatgtc
5101	cttccagggt	ctcagtgtcc	gcgtgagcgt	ggtctccgtc	acggtgaagg	ggtgggcccc
5161	agactataca	cttgcaaggg	tococttoao	actcatcctg	ctggtgctga	aacgggcacg
2101	gggctgcgcg	tacacataa	casastsacs	gttgaccatg	agetegtagt	tgagggcctc
222T	gtettegeee	racacarcaa	cyagatagea	getgaceaeg	agecegeage	ogggggggg
5281	ggcggcgtgg	cccttggcgc	ggagettgee	cttggaagag	egeeegeagg	cgggacagag
5341	gagggattgc	agggcgtaga	gcttgggtgc	gagaaagacg	gactcggggg	cgaaagcatc
5401	cactecacaa	tagacacaga	cggtctcgca	ctcgaccagc	caggtgagct	cgggctgctc
5461	agaatcaaaa	accapttttc	ccccattctt	tttgatgcgc	ttcttacctc	gcgtctccat
2#0T	gggccaaaa	accagettee	******	actatatata	teccentaga	conacttoat
552I	gagtetgtgt	eegegetegg	Lyacadacay	gctgtctgtg	ccccgcaga	cggacccgac
5581	gggcctgtcc	tgcaggggcg	teeegeggte	ctcctcgtag	agaaacccgg	accactctya
5641	gacgaaggcg	cgcgtccacg	ccaagacaaa	ggaggccacg	tgcgaggggt	agcggtcgtt
5701	gtccaccagg	gggtccacct	tttccacggt	atgcagacac	atgtccccct	cctccgcatc
5761	goodaodagg	attaacttat	aggtgtagge	cacgtgaccc	agaataccca	acqqqqqqt
2/0T	caagaaggcg	accygectyt	aggegeagge	-atatattaa	gggggggggg	ccacaaaaa
5821	ataaaagggg	gegggtetgt	getegteete	actctcttcc	geglegetge	t
5881	cagctgttgg	ggtaggtatt	ccctttcgag	agcgggcatg	accccggcac	teaggtigle
5941	agtttctaga	aacgaggagg	atttgatgtt	ggcttgccct	gccgcaatgc	tttttaggag
6001	actttcatco	atctggtcag	aaaagactat	ttttttattg	tcaagcttgg	tggcgaagga
6061	acceteaaaa	acattaneae	naancttooc	gatggatctc	atootctoat	ttttgtcacg
000T	gccaragagg	gegeeggaga	gaageeegge	gacggactco	toggeogee	cacacttaca
6121	gtcggctcgc	teettggeeg	cgatgttgag	ctggacatac	tegegegega	
6181	ttcggggaag	acggtggtgc	gctcgtcggg	cacgatectg	acgcgccagc	cgcggttatg
6241	cagggtgacc	agatccacgc	tggtggccac	ctcgccgcgc	aggggctcgt	tggtccagca
6301	ascacat cca	cccttacaca	agcagaacgg	gggcagcaca	tcaagcagat	gctcgtcagg
C2 C1	gaggagaaaa	togatgatas	agatroccor	acagagttcc	ttotcasaat	aatcgatttt
0207	ggggteegea	ccyacygcya	agatgeeegg	acagageeee	~~~~~~	actockooc
6421	tgaggatgca	tcatccaagg	ccatctgcca	ctcgcgggcg	geeagegete	getegtaggg
6481	gttgaggggc	ggaccccagg	gcatgggatg	cgtcagggcg	gaggcgtaca	tgccgcagat
6541	gtcgtagaca	tagatgggct	ccgagaggat	gccgatgtag	gtgggataac	agegeeeec
6601	gengatheth	acacacacat	antrataraa	ctcgtgcgag	ggggccaaga	aggcggggcc
0001	geggaegeeg	gegegeaege	**********	~22C2CC2+C	taacassas	taacatacaa
PPPT	gagarrggrg	egetgggget	gereggegeg	gaagacgatc	tygtgaaaga	cggcacgcga
6721	gttggaggag	atggtgggcc	gttggaagat	gttaaagtgg	gcatgaggca	gacgaaccga
6781	gtcgcggatg	aagtgcgcgt	aggagtcttg	cagcttggcg	acgagctcgg	cggtgacgag
6841	gacgtccatg	acacaataat	ccagcgtttc	gcggatgatg	tcataaccco	cctctccttt
C001	attataaast	2-2-2-3-049-	tracrocate	ctcctcgtca	teetteeant	actecedad
PAOT	CLUCUCAL	agetegege			townst-	±0000009949
6961	cgggaatcct	cgatcgtccg	cacggtaaga	gcccagcatg	Lagaaatggt	Leacygeett
7021	gtagggacag	cagcccttct	ccacggggag	ggcgtaagct	tgagcggcct	rgcggagcga
7081	aatatacatc	agggcgaagg	tatccctgac	catgactttc	aagaactggt	acttgaaatc
7141	casateatea		uctcccara	ctcgaaatcg	atacacttct	t.cgagagggg
1147		eage-cyclyt	artesta	727777777	antaccaca	
7201	gccaygcaga	ycyaaagtga	cyccattgaa	gagaatcttg	ttestes==	graryaaart
7261	gcgggtgatg	cggaaagggc	ccgggacgga	ggctcggttg	ttgatgacct	gggcggcgag

7221		tersarcert	tratottoto	cccgacgatg	tagagttcca	tgaatcgcgg
7201	gacgacttta	atatacaaca	actttttaag	ctcctcgtag	gtgaggtcct	cggggcaatg
7381	geggeetta	tactoraco	ccactcctg	gagatgtggg	ttggcttgca	tgaatgaagc
/441	cagteegtge	caccastas	agatetagaa	ctcgtcgcga	aagaggggga	actoctoocc
7501	ccagageteg	bbbbabaaa	tracreages	gaaagtaagg	gaateceact	cccagcgatc
/561	cacggccate		ratcacaaca.	gagggcgacc	agetetgggt	cccccaaaa
7621	ccagcgcaag	egeacggera	gategegage	cttgccgaag	gaccccatcc	aggtgtaggt
7681	tttcataacc	agcataaagg	ggacgagetg	cgtgcgagga	tgagagccga	ttgggaagaa
7741	ttctacatcg	taggtgacaa	agageegeee	cotottcato	tgatgaeagt	agaaatcccg
7801	ctggatttcc	tgccaccagt	cggacgagtg	gctgttgatg	ccacaatact	cocaococto
7861	ccggcgaacc	gagcactegt	getgatgett	gtaaaagcgt	ttgaggaga	acttcaggag
7921	cacgggctgt	acctcatcca	cgagatacac	agegegteec.	cectcecct	agaactecte
7981	tggcggccct	gacragragr	tttcatgttc	gcctgcgtgg	cacatetecc	cacaacaaaa
8041	gaggacggag	aggctgacga	geeegegegg	gagccaggtc	ataatataa	goggogggg
8101	gcggagagcg	aagacgaggg	egegeaging	ggagctgtcc	atgaracat	acttagasta
8161	gtccgggggc	agggttctga	ggttgacete	gtagaggcgg	tocacocatt	gcatgagaca
8221	cagatggtac	ttgatctcca	egggtgagtt	ggtggctgtg	agaagggatg	teacaracae
8281	gtagctgcgc	ggggccacga	ccgtgccgcg	gtgcgctttt	agaageggeg	cacatcaaca
8341	gctcccggcg	gcagcggcgg	tteeggeeee	gcgggcaggg	taacatacac	cacacacac
8401	tggcgctcgg	gcaggtcccg	grgergegee	ctgagagcgc	coaccccat	gacgacgogg
8461	cggttgacat	cctggatctg	cegeetetge	gtgaagacca	cagacacata	acccandate
8521	ctgaaagaca	gttcaacaga	atcaatctcg	gegteattga	acatosacto	ctccatctcc
8581	tcttgcacgt	cgcccgagtt	gteetggtag	gcgatctcgg	acatgaactg	attacanata
8641	tectectgga	gatcgccgcg	geeegegege	tccacggtgg	tagagagaga	actateaecc
8701	cgacccatga	gctgcgagaa	ggcgcccagg	ccgctctcat	caracttan	ctccacatac
8761	acgtccccgt	cggcgtcgcg	cgcgcgcatg	accacctgcg	cyayyccyay	actacacata
8821	cgcgtgaaga	cggcgtagtt	gcgcaggcgc	tggaagaggt	agittagggt	ggtggcgatg
8881	tgctcggtga	cgaagaagta	catgatccag	cggcgcaggg	geateteget	anactrosco
8941	atggcctcca	gcctttccat	ggcctcgtag	aaatccacag	cyaagiigaa	aaaccgggcg
9001	ttgcgggccg	agaccgtgag	ctcgtcctcc	aggageetga	tgagttegge	gatggtggtg
9061	cgcacctcgc	gctcgaaatc	cccgggggcc	tectectett	cetettette	catgacgacc
9121	tcttcttcta	tttcttcctc	tgggggcggt	ggtggtggcg	gggeeegaeg	acqueggega
9181	cgcaccggga	gacggtcgac	gaagcgctcg	atcatctccc	egeggeggeg	acycatygut
9241	tcggtgacgg	cgcgaccccg	ttcgcgagga	cgcagcgtga	agaegeegee	ggttatetet
9301	cggtaatggg	gcgggtcccc	gttgggcagc	gagagggcgc	tgacgatgca	thereers
9361	tgcggtgtag	gggacgtgag	cgcgtcgaga	tcgaccggat	cggagaatet	-t-cyayyaaa
9421	gcgtctagcc	aatcgcagtc	gcaaggtaag	ctcaaacacg	tagcagccct	gradacacta
9481	ttagaattgc	ggttgctgat	gatgtaattg	aagtaggcgt	ttttaaggeg	geggatggtg
9541	gcgaggagga	ccaggtcctt	gggtcccgct	tgctggatgc	gaageegete	ggecatgete
9601	caggcctggc	cctgacaccg	gctcaggttc	ttgtagtagt	catgcatgag	cctctcaatg
9661	tcatcactgg	cogaggegga	atcttccata	cgggtgaccc	cgacgcccct	gageggetge
9721	acgagcgcca	ggtcggcgac	gacgcgctcg	gcgaggatgg	cctgttgcac	gegggtgagg
9781	gtgtcctgga	agtcgtccat	gtcgacgaag	cggtggtagg	ccccggtgtt	gatggtgtag
9841	gtgcagttgg	ccatgagcga	ccagttgacg	gtctgcaggc	egggttgcac	gacctctgag
9901	tacctgagcc	gcgagaaggc	gcgcgagtcg	aagacatagt	cgttgcaggt	gegeaegayy
0061	tactootato	caactaggaa	atacaacaac	gactggcggt	agageggeea	gegergggrg
10021	accadededed	ccaaaaccaa	atcetegage	atgaggcggt	ggtagccgta	gaggtagtgg
10021	racatccagg	taataccaac	aacaataata	gaggcgcgcg	ggaactegeg	gacgcggttc
10141	cadatottoc	acaacaacaa	gaaatagtcc	atggtcggca	eggtetggee	ggtgagatgt
10201	acaceatcat	tgacgeteta	gaggcaaaaa	cgaaagcggt	tgagcgggct	ctteeteegt
10261	aggetageag	aacgcaaacg	gattagacca	catatatacc	ccggttcgag	tecectegaa
10321	teaggetaga	geogegaeta	acotootatt	ggcactcccg	tetegaeeeg	agecegatag
10381	ccaccaaaat	accoccocada	accetttttg	ccgaccgagg	ggagtegeta	gactigadag
10441	caaccaaaaa	ccccaccaaa	tagtggctcg	cqcccgtagt	ctggagaagc	tttgccaggg
10501	ttractcoco	gcagaacccg	attcacaaa	agccgcggcg	agcgggactt	ggtcaccccg
10561	ccmatttaaa	gacccacage	cacccgactt	ctccagttac	gggagegage	CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC
10621	ctttttacca	gatgeatece	atectacaco	aaatgcgtcc	Cacccccct	ccggcgacca
10691	CCCCCCCCCCC	gaccatagea	aacaccaaca	ctgtagcccc	gccacagcag	acagagatgg
10741	acttogaaga	gggcgaaggg	ctggcgagac	tgggggegee	gtccccggag	cgacaccccc
10901	acatacaact	gcagaaggag	atacacccaa	cgtacgtgcc	tgcgcagaac	ctgttcaggg
10061	accoractor	ggaggaggcc	gaggagatgc	gcgactgccg	ttttcgggcg	ggcagggagc
10921	Lacacaaaaa	cctggaccgc	cagcgcgtgc	tgcgcgacga	ggatttcgag	ccgaacgagc
1001	3-5-5-55			_		

FIG. 16A-3

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				L	assactaata	acancetaca
10981	agacggggat	cageceegeg	cgcgcgcacg	rggeggegge	caacciggig	acggeetacg
11041	agcagacggt	gaagcaggag	cgcaacttcc	aaaagagttt	caacaaccat	grgcgcacge
11101	taatcococo	caaaaaaata	accetagget	tgatgcacct	gtgggacctg	gcggaggcca
11161	tentacanaa	cccggacagc	aageetetga	caacacaact	attectagta	gtgcagcaca
11221	~~~~~~~	cgaggcgttc	addaaddcac	toctasacat	caccasaccc	gagggcgct
11221	gcagggacaa	cyaggegeee	agggaggege	gotagacat	acsaasacac	acctaacc
11281	ggctgctgga	gctgatcaac	atettgeaga	geategrage	gcaygagcgc	agcccgagcc
11341	tggccgagaa	ggtggcggct	atcaactact	cggtgctgag	cctgggcaag	ttttaegege
11401	gcaagattta	caagacgccg	tacgtgccca	tagacaagga	ggtgaagata	gacagctttt
11461	acatococat	ggcgctcaag	gtgctgacgc	tgagcgacga	cctgggcgtg	taccgcaacg
11521	accocateca	caaggccgtg	agegegagee	agcagcacaa	gctgagcgac	cgcgagctga
11501	tactacatet	gegeeggeg	ctaataaaaa	acaccaccaa	caataaaaa	tcctacttcg
11361		ggacctgcat	taaaaaaaaa	202002000	cttagaggcc	acctacaate
11641	acatgggggc	ggaccugcat	tygcageega	geeggegege	cccggaggcc	gaatactaac
11701	cagaggactt	ggatgaggat	gaggaagagg	aggaggatge	accegetgeg	gggtactgac
11761	gcctccgtga	tgtgttttta	gatgcagcaa	gccccggacc	ccgccataag	ggcggcgctg
11821	caaagccagc	cgtccggtct	agcatcggac	gactgggagg	ccgcgatgca	acgcatcatg
11881	gccctgacga	cccgcaaccc	cgagtccttt	agacaacagc	cgcaggccaa	cagactctcg
11941	gccattctgg	aggcggtggt	ccctctcgg	accaacccca	cgcacgagaa	ggtgctggcg
12001	atcotoaaco	cgctggcgga	gaagaaggcc	atccatccca	acgaggccgg	gctggtgtac
12001	accycyaacy	tggagcgcgt	addecate	aacagcacaa	acgtgcagtc	caacctggac
12001	aacyccctgc	Lygagegege	gggccgccac	accedence	accorttcaa	daacdaddac
12121	cggctggtga	cggacgtgcg	cgaggeegeg	gegeagegeg	ageggeeeaa	actacacaca
12181	ctgggctcgt	tggtggcgct	gaacgccttc	ctggcgacgc	ageeggegaa	
12241	gggcaggacg	attacaccaa	ctttatcagc	gegetgegge	tgatggtgac	cgaggtgccc
12301	cagagcgagg	tgtaccagtc	gggcccagac	tactttttcc	agacgagccg	gcagggcttg
12361	cagacggtga	acctaagcca	ggctttcaag	aatctgcgcg	ggctgtgggg	cgtgcaggcg
12421	cccatagaca	accggtcgac	ggtgagcagc	ttgctaacgc	ccaactcgcg	gctgctgctg
12421	atastasta	cgcccttcac	cuacaacaac	agcgtgaacc	gcaactcgta	cctgggccac
1240T	-tt	tttaccgcga	accestsacc	asaacacsaa	tanacaaaca	gaccttccag
12241	etgetgaege	Litadegega	ggccataggc	caggegeagg	cogacgagaa	dadadccacc
12601	gagatcacta	gcgtgagccg	cacacraaar	Cagaacgaca		
12661	ctgaacttct	tgctgacaaa	tagacagcag	aagattccgg	cgcagtacgc	getgteggee
12721	gaggaggagc	gcatcctgag	atatgtgcag	cagagcgtag	ggcttttcct	gatgcaggag
12781	ggggccaccc	ccagcgccgc	gctggacatg	accgcgcgca	acatggaacc	tagcatgtac
12841	gccgccaacc	ggccgttcat	caataagctg	atggactacc	tgcaccgcgc	ggctgccatg
12011	aactcggact	actttactaa	toctatacta	aacccccact	gactcccacc	gccggggttc
12061	tacceggace	agtacgacat	accessees	aacqatqqqt	tectatagaa	cgacgtggac
12301	cacacgggcg	agtacgacat	geeegaeeee	accourages	caatacacac	acccacasac
13021	agcgcggtgt	tctccccgac	cccgcaaaag	cgccaggagg	cygtacycac	attacaaaa
13081	gagggcgcgg	tgggtcggag	ccccttcct	agcttaggga	guuguatag	-t
13141	tcggtgaaca	gcggcagggt	gagccggccg	cgcttgctgg	gcgaggacga	gtacctgaac
13201	gactcgctgc	tgcagccgcc	gcgggtcaag	aacgccatgg	ccaataacgg	gatagagagt
13261	ctaataaaca	aactgaaccg	ctggaagacc	tacgctcagg	accataggga	tgcgcccgcg
13321	ccacaacaac	agcgccacga	ccaacaacaa	aacctaatgt	gggacgacga	ggactcggcc
13381	gacgatagea	gcgtgttgga	cttgggggg	agcagtagag	ccaacccgtt	cgcgcatctg
12201	gacgacagea	tggggcgacg	detatttta	atraaataaa	actcaccaag	gccatagcgt
T344T	cageceagae	cggggcgacg	gatgettega	acguactat	cttcctctcc	tecteceted
13501	gegttetett	ccttgttaga	gatgaggege	geggege	tteteetee	
13561	tacgagagcg	tgatggcgca	ggcaaccctg	gaggttccgt	Ligigactica	geggeatate
13621	gctcctacgg	agggcagaaa	cagcattcgt	tactcggaac	tggctccgca	gtacgacacc
13681	actcgcgtgt	acttggtgga	caacaagtcg	gcggacatcg	cttccctgaa	ctaccaaaac
13741	gaccacagca	acttcctgac	cacggtggtg	cagaacaacg	atttcacccc	cgccgaggcc
13801	agcacgcaga	cgataaattt	tgacgagggg	tcacaatagg	gcggtgattt	gaagaccatt
13861	ctocacacca	acatgcccaa	tataaacaaa	tacatottca	ccagcaagtt	taaggcgcgg
12001	etesteetee	ctaggaaggt	agtgaatgag	aatgatagga	gcaaggatga	gttaaaatat
T327T	gracegrag	ccaygaaggc	ggcagaccag	ancyatty	araccatrac	catagacctg
13381	gagtggtttg	agtttaccct	geeegaggge	additteeg	aguccacgac	catagacceg
14041	atgaacaacg	ccatcttgga	aaactacttg	caagtggggc	ggcaaaacgg	cgtgctggag
14101	agcgatatcg	gagtcaagtt	tgacagcagg	aatttcaagc	tgggctggga	cccggtaacc
14161	aagctggtga	tgcctggggt	ctacacctac	gaggccttcc	acccggacgt	tgtgctgctg
14221	ccaaactaca	gggtggactt	caccgagagc	cgcctgagca	acctcctggg	cattcgcaag
14281	aagcaacctt	tccaagaggg	cttcaggate	atgtatgagg	atctcgaggg	tggtaacatc
1 42 41	coccccata	tggatgtcaa	dcaatattta	gatagtaaaa	agaagettga	ggaggcaaca
1474T		ccagggctgc	tageactet	adaddadada	gtcatattca	aanagetoto
14401	cagaatgcaa	ccayggetge	Lygayatatt	ayayyayaca		2242242
14461	gaacaagcgg	ctgaaaagga	cctggtcatt	gtaccagtaa	cacaayatga	aaytaayaya
14521	agctataatg	tcatagatgg	cacccatgac	accetetace	gaageeggta	cctgtcctat
14581	acctacgggg	accccgagaa	gggggtgcag	tcgtggacgc	tgctcaccac	cccggacgtc

FIG. 16A-4

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14641	acctacaaca	consocasot	ctactootco	ctoccooacc	tcatgcaaga	ccccatcacc
14701	ttaagatata	ccaccaat	caccaactac	cccataatta	gcgccgagct	catoccette
14761	LLCCGCLCLA	cctagtaagt	cagcaaccac	atatactaca	agctcatccg	carctacacc
14/61	cgcgccaaga	gettttaeaa	b	gtttactett	tactatacas	tagacacaca
14821	teceteacee	acgtetteaa	eegetteeee	gacaaccaga	tectetgeeg	coccecegeg
14881	cccaccatca	ccacggtcag	tgaaaacgtg	cetgetetea	cagatcacgg	gacycraccy
14941	ctgcgcagca	gtatccgcgg	agtccagcga	gtgaccgtca	ctgacgcccg	tegeegeace
15001	tgtccctacg	tctacaaggc	cctgggcata	gtcgcgccgc	gcgtgctttc	cagtcgcacc
15061	ttctaaaaaa	tgtctattct	catctcgccc	agcaataaca	ccggctgggg	tcttactagg
15121	cccagcacca	tgtacggagg	agccaagaag	cgctcccagc	agcaccccgt	ccgcgtccgc
15181	ggccacttcc	gcgctccctg	gggcgcttac	aagcgcgggc	ggacttctac	cgccgccgtg
15241	cgcaccaccg	tegacgacgt	catcgactcg	gtggtcgccg	acgcgcgcaa	ctataccccc
15301	acccctcca	ccatagacac	ggtcatcgac	agcatagtag	ccgacgcgcg	cgactatgcc
15361	acacccccca	accaacaaca	acquatcqcc	aggcgccacc	ggagtacgcc	cgccatgcgc
15/21	agacgcaaga	ctctactaca	ccgcgccaga	cacacaaacc	gccgggccat	gatgcgagcc
15/01	geegeeeggg	ccaccactac	accccccca	ggcaggactc	gcagacgagc	aaccaccacc
1040T	gegegeegeg	costttctac	catraccara	cccadacaca	gaaacgtgta	ctaggtagge
10041	getgeegegg	ccattttttag	catgaccaga	cccaggegeg	ctcctcgtcc	ctgatctaat
12001	gactccgtca	egggegtgeg	egracedra.	cgcacccgcc	aatraarrar	cagatoctac
15661	gcttgtgtcc	tcccccgcaa	gegaegaege	caaaycycaa	aatcaaggag	gagatyctcc
15721	aggtcgtcgc	cccggagatt	tacggaccac	cccaggcgga	ccagaaaccc	cycaaaatca
15781	agcgggttaa	aaaaaaggat	gaggtggacg	agggggcagt	agagtttgtg	egegagiteg
15841	ctccgcggcg	gcgcgtaaat	tggaaggggc	gcagggtgca	gcgcgtgttg	cggcccggca
15901	cggcggtggt	gtttacgccc	ggcgagcggt	cctcggtcag	gagcaagcgt	agctatgacg
15961	aggtgtacgg	cgacgacgac	atcctggacc	aggcggcgga	gcgggcgggc	gagttcgcct
16021	acgggaagcg	gtcgcgcgaa	gaggagctga	tctcgttgcc	gctggacgag	agcaacccca
16081	cocctaocct	gaagcccgtg	accetgeage	aggtgctgcc	ccaagcagtg	ctgctgccga
16141	accacaaaat	caagcgcgag	ggcgagaata	tgtacccgac	catgcagatc	atggtgccca
16201	acccccaca	cotogaagaa	gtgctggaca	ccgtgaaaat	ggatgtggag	cccgaggtca
16261	agagaagagag	catcaagcag	ataacaccaa	acctagacat	gcagaccgtg	gacattcaga
16331	tecceacea	categatgtt	gacaaaaaac	cctcgaccag	catcgaggtg	cagaccgacc
16301	catacataca	acctccacc	actaccatet	ccacttctac	cgccgccacg	octaccoaoc
10301	change	agccccacc	geegeegee	acconctoat	gcccaactac	gtattgcatc
10441	ccccagaag	gcgaagatgg	ggccccgcca	accegecegae	ctacgccagc	cacsaacacc
T020T	CTTCCattat	ecegacyccy	ggetategeg	gcacccggca	tetegecage	accordatec
16561	cagccagcaa	acgccgccgc	egeacegeea	ceegeegeeg	tetggeecee	tagagaga
16621	gccgcgtaac	cacgcgccgg	ggeegetege	tegttetgee	caccgtgcgc	Laccacccca
16681	gcatccttta	atccgtgtgc	tgtgatactg	ttgcagagag	atggctctca	cttgeegeet
16741	gcgcatcccc	gtcccgaatt	accgaggaag	atcccgccgc	aggagaggca	tggcaggcag
16801	cggcctcaac	cgccgccggc	ggcgggccat	gegeaggege	ctgagtggcg	gctttctgcc
16861	cgcgctcatc	cccataatcg	cggcggccat	cggcacgatc	ccgggcatag	cttccgttgc
16921	actacagaca	tcgcagcgcc	gttgatgtgc	gaataaagcc	tctttagact	ctgacacacc
16981	taatectata	tatttttaga	atggaagaca	tcaattttgc	gtccctggct	ccgcggcacg
17041	acacacaacc	attcatgggc	acctggaacg	agatcggcac	cagccagctg	aacgggggcg
17101	ccttcaattq	gagcagtgtc	tagaacaaac	ttaaaaattt	cggctcgacg	ctccggacct
17161	atoronacaa	ggcctggaat	agtagcacgg	ggcagttgtt	aagggaaaag	ctcaaagacc
17721	acgggaacaa	ggeoogganta	ataascaacc	tagecteggg	cattaacggg	gtggtggaca
17201	tomassaga	gaagaaggag	cacaaataa	acadecdect	ggacccgcgg	ccacccacaa
17241	Laguadacca	ggccgcgcag	actestes	caccasaaa	cgagaagcgg	ccacaaccca
1/341	tggtggagat	gyaayatyta	accecege	cycccaayyy	atacaaaaa	ccacaacca
17401	acgcggagga	gacgatecty	caggiggacg	ageegeeeee	gtacgaggag	gccgccaagg
17461	ccggcatgcc	caccacgcgt	accategege	caccggccac	tggtgtaatg	adaceegeed
17521	cccttgacct	gcctccgcca	cccacgcccg	ctccaccgaa	ggcagctccg	grigingeage
17581	cccctcctgt	ggcgaccgcc	gtgcgccgcg	teceegeeeg	ccgccaggcc	cagaactggc
17641	agagcacgct	gcacagtatc	gtgggcctgg	gagtgaaaag	tctgaagcgc	cgccgatgct
17701	attgagagag	aggaaagagg	acactaaagg	gagagcttaa	cttgtatgtg	ccttaccgcc
17761	agagaacgcg	cgaagatggc	taccccctcg	atgatgccgc	agtgggcgta	catgcacatc
17821	accadacaga	acgcctcgga	gtacctgagc	ccgggtctgg	tgcagtttgc	ccgcgccacc
17881	gacacgtact	tcagcctggg	caacaagttt	aggaacccca	cggtggctcc	cacccacgat
17941	ataaccacaa	accogtccca	gcgtctgacg	ctgcgctttg	tgcccgtgga	tcgcgaggac
18001	accacatact	catacaaaac	gcgcttcact	ctaaccataa	gcgacaaccg	ggtgctagac
18061	atorceares	cttactttca	cateegegge	atcatagaca	gcggtcccag	cttcaaaccc
10101	tactororo	caacttacaa	cadectage	CCCSSSCOO	ccccaactc	tagtcagtgg
10101	annanas-	angetacea	taccaatese	aarraaactr	acacatttgg	actaccccc
TOTOT	yaacaagcta	aayetaccaa	antnassnot	cttcasatta	gaactgatga	aacta=ccs=
19741	argggcggag	aagacattac	ugrgaaaygt	ccccaaaccy	Jacobyacya	uactaayyaa

FIG. 16A-5

			•			_
18301	gatggagagg	atgaaatttt	tgcagatcaa	acattccagc	cagaacctca	agtgggagaa
18361	cagaactggc	aagaaacott	tottttctat	ggaggcagag	ctcttaagaa	agaaaccaaa
18/21	atgaagccat	attataactc	ttatgcgaga	cccacaaato	aaaagggagg	acaggetaaa
10422	tttacacttg	atrassaarr	tranccaacc	aaaattcctg	atattacaat	ggatttcttt
10541	gatagtccac	acyaaaaaagg	atcagecuate	actastasco	caratattat	catotatoca
18541	gatagreeae	aagatgatac	a caygegea	actaataayt	cagacaccyc	caegeaegea
18601	gaaaatgtaa	atttagaagc	teetgacaca	cargragere	acaaaccayy	caaagacgac
18661	tctagttctt	ccgctaacct	cacacaacag	gccatgccta	acagaccgaa	ctacateggg
18721	ttcagagaca	actttgtggg	tcttatgtac	tacaatagta	ctggcaacat	gggtgtgctg
18783	gctggtcagg	cctctcagtt	gaatgctgtg	gtcgacttgc	aagacagaaa	caccgagctg
18841	tcttaccagc	tattgctaga	ttctctgggt	gacagaacca	gatactttag	catgtggaat
18901	tctgcagtgg	acagctatga	ccccgatgtc	aggatcattg	agaatcacgg	tgtggaagat
18961	gaacttccaa	actattgctt	cccactgaat	ggcagtggtt	ctaacagcac	atacaaaggt
19021	gttaaagctg	gaactggaaa	caattgggat	gacgatgaaa	atgttgcaag	acaaaatcag
19081	attggcactg	gcaacctgtt	cgccatggag	atcaacctcc	aggccaacct	atggaagagt
19141	tttctgtact	cgaacgtggc	cctgtacctg	cccgactcct	acaagtacac	gccggccaac
19201	gtcacgctgc	ccaccaacac	caacacctac	gactacatga	acaaccacat	ggtagccccc
19261	tegetggtgg	acacctacat	caacattggc	acccactaat	cactagaccc	catogacaat
10201	gtcaatccct	tcaaccacca	ccacaecaca	gacctacact	accostscat	actectagae
10201	. aacggccgct	acataccatt	ccacatccaa	gtgccgcge	agttetttge	catcaagaac
	ctgcttctgc					
1944	atcctgcaga	-tto-ctggttc	ccacacctac	gagaggaacc	coccetecat	ccacttcasc
19501	. accetgeaga	gricectegg	caacgacctg	cycytcyacy	gegeeeege	cagactagae
1956	. agcgtcaacc	tetaegecae	ettetteeee	acggegeaca	acaccyccic	caccetggaa
1962	. gccatgctgc	gcaacgacac	caacgaccag	tectteaacg	actacetete	ggccgccaac
1968	. atgctctacc	ccatcccggc	caaggccacc	aacgtgccca	tetecatece	ctcgcgcaac
19743	. tgggccgcct	teegeggetg	gagtttcacc	cggctcaaga	ccaaggaaac	recereceré
19801	ggctcgggtt	tcgaccccta	ctttgtctac	tegggeteca	tcccctacct	cgacgggacc
19861	ttctacctca	accacacctt	caagaaggtc	tccatcatgt	tcgactcctc	ggtcagctgg
1992	. cccggcaacg	accggctgct	cacgccgaac	gagttcgaga	tcaagcgcag	cgttgacggg
19981	gagggctaca	acgtggccca	atgcaacatg	accaaggact	ggttcctcgt	ccagatgctc
20043	. tcccactaca	acatcggcta	ccagggcttc	cacgtgcccg	agggctacaa	ggaccgcatg
20101	tactccttct	tccgcaactt	ccagcccatg	agcaggcagg	tggtcgatga	gatcaactac
20161	. aaggactaca	aggccgtcac	cctacccttc	cagcacaaca	actegggett	caccggctac
20221	. cttgcgccca	ccatacacca	gggggagccc	taccccccca	acttccccta	cccactcatc
20281	. ggctccaccg	cagttccctc	cotcacccao	aaaaagttcc	tctgcgacag	ggtcatgtgg
2020	. cgcatcccat	tetecareaa	ctttatgtcc	atagacaccc	tcaccgacct	gogtcagaac
	. atgctctatg					
	. gageceacce					
	. caccgcggcg					
2058.	. acttaagcat	gageggetee	agegaacaag	agetegege	teratterst	gaccigggat
2064.	. gcgggcccta	ctttttggga	acccacgaca	agegetteee	tggçtteett	geeggegaca
2070	. agctggcctg	cgccatcgtc	aacacggccg	gccgcgagac	cggaggcgcg	cactggeteg
2076	. cctttggctg	gaatccgcgc	tcgcgcacct	gctacatgtt	cgaccccttt	gggttctcgg
20823	. accgccggct	caagcagatt	tacagcttcg	agtacgaggc	catgctgcgc	cgaagcgcgc
2088:	. ttgcctcctc	gcccgaccgc	tgtctcagcc	tcgagcagtc	cacccagacc	gtgcaggggc
20943	. ccgactccgc	cgcctgcgga	cttttttgtt	gcatgttttt	gcatgccttc	gtgcactggc
21003	. ccgaccgacc	catggacgga	aaccccacca	tgaacttgct	gacgggggtg	ccaaacggca
21063	tgctacaatc	gccacaggtg	ctgcccaccc	tcaggcgcaa	ccaggaggag	ctctaccgct
2112	. tcctcgcgcg	ccactcccct	tactttcgat	cccaccgcgc	cgccatcgaa	aacgccaccg
2118	cttttgataa	aatgaaacaa	ctgcgtgtat	ctcaataaac	agcactttat	tttacatgca
2124	ctġgagtata	tocaagttat	ttaaaagtcg	aaggggttct	cgcgctcgtc	gttgtgcgcc
2130	gcgctgggga	gggccacgtt	acaatactaa	tacttgggaa	gccacttgaa	ctcggggatc
2136	accagtttgg	gcactggggt	ctcggggaag	atctcactca	acatococco	gctcatctgc
2142	agggcgccca	dratuterda	accadadate	ttgaaatcac	aattoggggg	gatactetac
2110	gegegegagt	tacaateceg	addattacea	cactoraaca	ccattagact	gaaateatta
2140	. gegegegagt Lacactggcaa	garagetatat	gyggccgcag	tactygaaca	coaccagaca	agg caccec
2134.	acactggcaa	gcacgctctt	gregergate	cyaccettgt	aggicette	otana-atta
2160	aggccgaacg	gggtcatctt	gcacayetgg	cygcccagga	agggcacgct	cryaggerig
2T00	tggttacact	cgcagtgcac	gggcatcagc	atcatccccg	egeegegetg	catatteggg
2172	tagagggcct	tgacgaaggc	cgtgatctgc	ttgaaagctt	gctgggcctt	agccccctcg
2178	ctgaaaaaca	ggccgcagct	cttcccgcta	aactggttat	tecegeacee	ggcatcatgc
2184	l acgcagcagc	gcgcgtcatg	gctggtcagt	tgcaccacgc	tacgtcccca	geggttetgg
2190	l gtcaccttgg	ccttgctggg	ctgctccttc	aacgcgcgct	gcccgttctc	gctggtcaca

FIG. 16A-6

21961	tccatctcca	ccacgtggtc	cttgtggatc	atcaccgtcc	catgcagaca	cttgagctga
22021	ccctcgacat	cocaocaocc	atgatcccac	agggcgcagc	cggtgcactc	ccagttetta
22001	tacacattca	cactataact	gaagatgtaa	ccttqcaaca	ggcgacccat	gaeggugeta
221/1	satactttct	gggtggtgaa	gatcaattac	agaccgcggg	cctcctcgtt	catccaggic
22201	torcacatet	tttggaagat	ctcaatctac	tegggeatga	gcttgtaagc	accycycagy
22261	ceactateas	cacaataaca	ttccatcagc	acgttcatgg	tatecatgee	Ctttttccag
22221	macmamacca	gagggagact	cagggggttg	cccaccttca	ggacaccggg	ggtegeagge
22321	tegacgatge	attttcatc	cttaccttcc	ttcaacagaa	ccggaggctg	gctgaatccc
2230I	actcccacga	ttaccacatc	ttcctaagac	atctcttcgt	cagaatctac	cttggtcaca
77441	tgcttggtct	ttatagetta	cttcttttt	ggagggetgt	ccacaaaaac	cacgtcctcc
ZZ501	tcggaagacc	ccccggcccg	cccctcatac	tttcggcgct	tagtagacag	aggaggtggt
22561	ggcggcgagg	eggageceae	stastacaac	gratagraca	ccaeccata	accccadaac
22621	ggcggcgagg	ggeteetete	enge recegge	acatactasa	taccaccaac	cattotttcc
22681	ggagtggcct	ctegeteeat	gaaceggege	acgeeeegae	aggaggaggt	aaccacccac
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35101	ccctaaattc	aaaagctcat	ttgcatgtta	acttttgttt	actttgtggg	gtatattatt
35161	gatgatc					
SEO II	NO: 5		•			

Grp	Vaccine	Monkey	P	re	W	k 4		K 8	Wk	12
	at Wk 0, Wk 4	ID	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Gag
1	Ad24AE 1gogAOrf6Ad5Orf6	00C072	3	4	4	381	3	150	3	68
	10^11 vp	00C178	3	3	1 1	559	1	743	0	635
		00C222	0	3	1 1	369	וו	753	0	670
		00D011	1 1	9	9	211	4	273	0	520
. •		00D023	0	6	0	295	1	459	1	368
		00D031	15	5	10	103	1	101	1	40
2	Ad24AE 1gogAOrf6Ad5Orf6	99C168	4	6	0	118	5	241	3	209
-	10^10 VP	99C170	10	5	5	241	3	141	3	103
-	10 10 1	99C173	1	.3	0	23	0	14	0	21
3	Ad24ΔE1gcgΔE4Ad5Orf6	99C154	0	3	0	93	0	60	1	53
•	10^10 vp	99C158	1	0	1	141	0	101	1 1	120
	, , , , , <sub>F</sub>	99C177	Ð	0	0	45	0	39	٥	79
4	MRKAd5-HIVgag	00C018	1	5	13	1025	0	824	3	753
•	10^11 Vp	00C034	0	4	5	219	5	404	0	491
	•	00C058	4	4	3	1086	0	440	0	439
5	MRKAd5-HIVgag	99C218	0	. 3 .	5	2500	0	1580	10	1655
٠.	10^10 VD	99C227	6	1	4	529	5	365	5	1004
	/ <del>-</del>	99D185	ND	ND	0	425	0	310	0	271

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Vaccine	Monkey	Gag-Specific (Wk 1	
at Wk 0, Wk 4	ID	%CD4	%CD8
Ad24ΔE 1 gogΔOrf6Ad5Orf6	00C072	0.02	0.02
10^11 vp	00C178	0.05	0.38
•	00C222	0.02	0.40
	00D011	0.02	0.27
	00D023	0.01	0.11
	00D031	0.01	0.01
MRKAd5-HIVgag	00C018	0.05	0.41
10^11 vp	00C034	0.06	0.18
	00C058	0.02	0.28

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Grp	Vaccine	Monkey	Wk 4	WK 8
•	at Wk 0, Wk 4	ID		
1	Ad24AE 1 gag\Orf6Ad5Orf6	00C072	<10	77
	10^11 vp	00C178	<10	26
		00C222	<10	423
		00D011	<10	98
	•	00D023	<10	<10
		00D031	<10	<10
2	Ad24AE 1 gog AOrf6Ad5Orf6	99C168	<10	<10
	10^10 vp	99C170	<10	<10
		99C173	<10	<10
3	Ad24ΔE1gcgΔE4Ad5Orf6	99C154	<10	<10
	10^10 vp	99C158	<10	<10
	•	99C177	<10	<10
4	MRKAd5-HIVgag	00C018	34	1017
	10^11 vp	00C034	14	423
		00C058	46	934
5	MRKAd5-HIVgag	99C218	20	99
	10^10 vp	99C227	40	767
		99D185	17	342

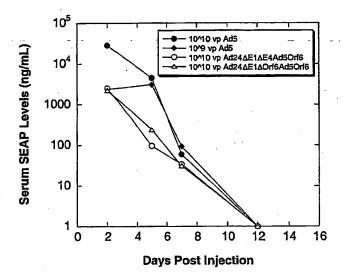


FIG. 20

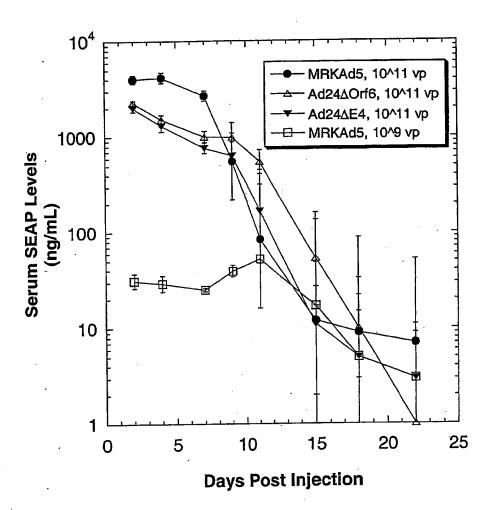
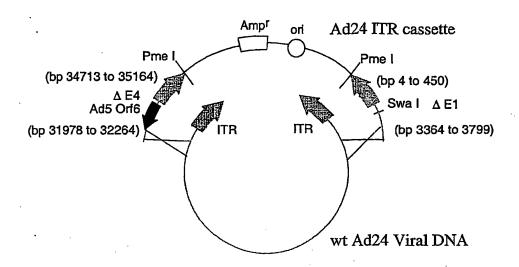


FIG. 21

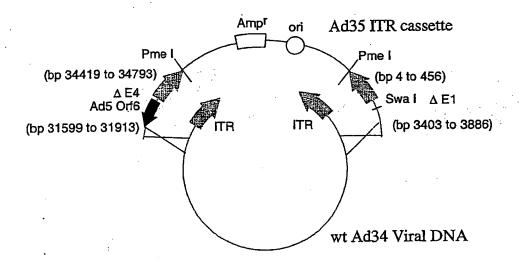
40/59.

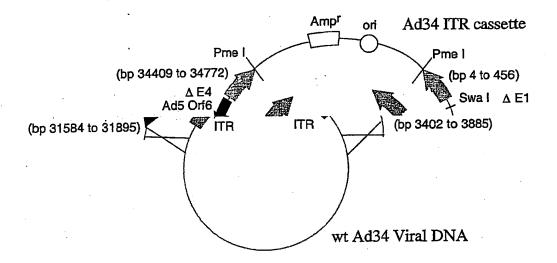


Animai	Prime (Wk 0, 4, 26)	Boost (Wk 56)	Pre		Prime <sup>b</sup>		Pre-Boost®		Post-Boost <sup>d</sup>	
Anımaı	Prime (WK U, 4, 20)	5555 (777 55)	Mock*	Gag*	Mock	Gag	Mack	Gag	Mock	Gag
Manhau 4	10 <sup>8</sup> vp MRKAd5-gag	1011 vp Ad24ΔE1gagΔOrf6Ad5Orf6	18	16	1	244	3	74	3	1235
Monkey 1	10 vp MRKAd5-gag	10 <sup>11</sup> vp Ad24AE1gagAOrf6Ad5Orf6	10	9	4	83	0	18	0	856
Monkey 2	10° vp MRKAd6-gag	1011 vp Ad24AE1gagAOrf6Ad5Orf6	l 1	1	۱ ،	219	9	69	0	703
Monkey 3 Monkey 4	10 vp MRKAd6-gag	10 <sup>11</sup> vp Ad24ΔE1gagΔOrf6Ad5Orf6	1	1	3	59	1	20	0	419
Mankey 5	none	10 <sup>11</sup> vp Ad24ΔE1gagΔOrf6Ad5Orf6	3	4	ND	ND	ND	ND	4	558
Monkey 6	nane	1011 vp Ad24AE1pagAOrf6Ad5Orf6	0	3	ND	ND	ND	ND	1	295
Monkey 7	none	1011 vp Ad24AE1gagAOrf6Ad5Orf6	1	9	ND	ND	ND	ND	8	103
	none	1011 vp Ad24AE1gagAOrf6Ad5Orf6	3	3	ND	ND	ND	ND	1	381
Monkey 8	none	1011 VD Ad24AE1gagAOrf6Ad5Orf6	0	6	ND	ND	ND	ND	0	369
Monkey 9 Monkey 10	none	1011 vp Ad24AE1pagAOriBAd5Ori6	15	5	ND	ND	ND	ND	10	211

Animal	Prime (Wk 0, 4, 26)	Boost (Wk 56)	Gag-Specific	<u> Cells (Wk 60)</u>
Aillinai			%CD4	%CD8
Monkey 1	10 <sup>9</sup> vp MRKAd5-gag	10 <sup>11</sup> vp Ad24∆E1gag∆Orf6Ad5Orf6	0.06	0.37
Monkey 2	107 vp MRKAd5-gag	10 <sup>11</sup> vp Ad24∆E1gag∆Orf6Ad5Orf6	0.01	0.56
Monkey 3	109 vp MRKAd6-gag	10 <sup>11</sup> vp Ad24∆E1gag∆Orf6Ad5Orf6	0.07	0.06
Monkey 4	107 vp MRKAd6-gag	10 <sup>11</sup> vp Ad24∆E1gag∆Orf6Ad5Orf6	0.04	0.20

Animai	Prime (Wk 0, 4)	Boost (Wk 24)	P	re	Prin	neb	Pre-B	oost*	Post-F	3oost <sup>d</sup>
·	, (, .,	, ,	Mock	Gag*	Mock	Gag	Mock	Gag	Mock	Gag
Monkey 11	1011 vp Ad244E1gagAOrf6Ad5Orf8	107 vp MRKAd5-gag	3	4	3	150	4	28	0	188
Monkey 12	1011 vp Ad24AE1gagAOrf6Ad5Orf6	107 vp MRKAd5-gag	0	3	1	753	4	554	٥	1029
Mankey 13	10 <sup>11</sup> vp Ad24∆E1gag∆Orl6Ad5Orl6	10 <sup>7</sup> vp MRKAd5-gag	1	9	4	273	0	370	0	1520
Monkey 14	none	10 <sup>7</sup> vp MRKAd5-gag	0	0	ND°	ND	ND	ИD	4	94
Mankey 15	none	107 vp MRKAd5-gag	0	0	ND ·	ND	ND -	ND	1	168
Monkey 16	none	107 vp MRKAd5-gag	8	3	ND	ND	ND_	ND	. 8	149





1	catcatcaat	aatatacctt	atagatggaa	tggtgccaat	atgtaaatga	ggtgatttta
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	cgtgggaaaa					
	acgcataaaa					
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	tgaggaagtg					
	ccaggtagac					
	ccgcgtaccg					
	tatacctcag					
	tetgegeegg					
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	agaggtagag					
	tatgctttta					
	tccaggggtg					
	ggactgtgat					
	aaaggagcag					
	tcagttggat					
1081	aaatactgga	gtaaaggaac	tgttatgttc	gctttgttat	atgagagcgc	actgccactt
	tatttacagt					
1201	attgagtggg	agttttgtgc	ttcttattat	aggtectgtg	tctgatgctg	atgagtcacc
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	taagagggag					
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2461	acagtataag	attactacac	ccattaatat	ccacactact	tattacatat	ctoreastor
2521	ggctgaggtg	gtaatagata	ctcaagacaa	ggcagttatt	agatgctgca	toatocatat
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3541	tcccatcctg	ggcaggagtt	catcadaata	ttatgggatg	tactgtggat	aggagaccca
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3661	cagctgcagc	cgccgccgcc	gcctctgttg	ccgctaacac	tgtgcttgga	atgggttact
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6661	geceeetet	catacttcct	cocacataot	catatacttc	atotoatooc	octaocaacc
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0/21	ccggacccaa	arragracas			gacaatetyy	tanaataa
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7001	agenttacae	gagttatata	trataataa	gctgtacctg	actteectta	acqaqaaatt
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0221	gagteetgag	acgccgcga	cccaggccag			h
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30601	aatgaattaa	aaaatgatta	ataaaaaatc	actiactiga	aatcagcaat	had a comme
30661	ttgaaatttt	ctcccagcag	cacctcactt	CCCTCTTCCC	aactctggta	LICTABACCC
30721	cattcaacaa	catactttct	ccatacttta	aaggggatgt	caaattttag	ctcctctcct
30781	gtaccacaa	tcttcatotc	tttcttccca	gatgaccaag	agagtccggc	tcagtgactc
30841	cttcaaccct	atctacccct	atgaagatga	aagcacctcc	caacacccct	ttataaaccc
20001	accedacee	tacacassta	cottoscoc	Tagadocada	ggagttctta	ctttassato
20201	ayyyttätt	Lucudadaug	gullicatata	tatagagaga	2244	coccatate
30961	tttaacccca	ctaacaacca	caggcggatc	cclacageta	aaagtgggag	ggggacttac
31021	agtggatgac	actgatggta	ccttacaaga	aaacatacgt	gctacagcac	ccattactaa
31081	aaataatcac	tctgtagaac	tatccattgg	aaatggatta	gaaactcaaa	acaataaact
31141	atgtgccaaa	ttgggaaatg	ggttaaaatt	taacaacggt	gacatttgta	taaaggatag

21201	tattaacacc	ttatagagta	raataaaccc	tocacctaac	tgtcaaattg	togaaaacac
31201	tactaacact	ccacggaccg	ttactttact	attactaaaa	aacggagggc	ttottaatoo
21701	taatacaaat	gatggtaaac	teaccitage	tetesseess	atottopoo	assauscauc
31321	ctacgtgtct	ctagttggtg	tattagatat	tytyaaccaa	atgttcacac	addagacago
31381	aaacatccaa	ttaagattat	attttgactc	ttctggaaat	ctattaactg	algaallaya
31441	cttaaaaatt	ccacttaaaa	ataaatcttc	tacagegace	agtgaaactg	tagccagcag
31501	caaagccttt	atgccaagta	ctacagctta	tcccttcaac	accactacta	gggatagtga
31561	aaactacatt	catogaatat	ottactacat	gactagttat	gatagaagtc	tatttccctt
31621	gaacatttct	ataatoctaa	acagccgtat	gatttcttcc	aatgttgcct	atgccataca
31681	atttgaatgg	aatctaaatq	caagtgaatc	tccagaaagc	aacatagcta	cgctgaccac
217/1	atcondettt	ttcttttctt	acattacada	эдасдасаас	taaaataaag	tttaagtgtt
31/41	tttattta	atananan	togactage	attttacctc	caccttccca	tttgacagaa
3190T	tttatttaaa	tetacaaaa	cogagiagit	accetttana	taccattaga	gatagacatt
31861	tacaccaatc	Ecceccacy	cacayettta	-accertings	atctactaga	antmatamat
31921	gttttagatt	ccacattcca	aacagtttca	gagcgagcca	atctggggtc	agryaragar
31981	aaaaatccat	cgcgatagtc	ttttaaagcg	ctttcacagt	ccaactgctg	cygacycyaa
32041	tccggagtct	ggatcacggt	catctggaag	aagaacgatg	ggaatcataa	teegaaaaeg
32101	gtatcggacg	attgtgtctc	atcaaaccca	caagcagccg	ctgtctgcgt	cgctccgtgc
32161	aactoctott	tatoggatca	gggtccacag	tgtcctgaag	catgatttta	atagccctta
32221	acatcaactt	tctggtgcga	tgcgcgcagc	aacgcattct	gatttcactc	aaatctttgc
32281	agtaggtaga	acacattatt	acaatattqt	ttaataaacc	ataattaaaa	gcgctccagc
32341	casaactcat	atctgatata	atcocccto	catgaccatc	ataccaaagt	ttaatataaa
22/101	ttaaatracr	ttccctcaaa	aacacactac	ccacatacat	gatctctttt	ggcatgtgca
22461	tottaaacgacg	ctctctctaa	catogaceac	attaattaat	catgcaaccc	aatataacct
32401	tattaataat	cogcocgcac	acceptance	carccatrica	ttgaagtgaa	ccctcctcat
32321	teeggaacca	cactyccaac	accyciccic	cagccacgcu	cacttcacaa	trassastat
32581	tacaatgaca	atgaagaacc	caattette	gaccycgaac	cacttgagaa	anatoataa
32641	ctatagtggc	acaacataga	cataaatgca	tgcatettet	cataattttt	abcccccag
32701	gatttagaaa	catatcccag	ggaataggaa	gctcttgcag	aacagtaaag	etggeagaac
32761	aaggaagacc	acgaacacaa	cttacactat	gcatagtcat	agtatcacaa	tctggcaaca
32821	acaaataatc	ttcagtcata	gaagctcggg	tttcattttc	ctcacaacgt	ggtaactggg
32881	ctctggtgta	agggtgatgt	ctagcacata	atgtcgagcg	tgcgcgcaac	cttgtcataa
32941	togagttgct	tcctgacatt	ctcgtatttt	gtatagcaaa	acgcggccct	ggcagaacac
33001	actettette	gccttctatc	ctgccgctta	acatattcca	tgtgatagtt	caagtacagc
33061	cacactetta	anttontcaa	aagaatgctg	gcttcagttg	taatcaaaac	tccatcgcat
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33121	ctaattgett	tttaaaaaaa	accordaged	goataggaa	gaagaaccat	gttaattttt
33191	cuggatugeg	ttttaagtag	gagagagag	ggaagagacg	caratrorat	ctctcacccc
33241	attccaaacg	atetegeagt	acticaaatt	grayarcycy	cagatggcat	aggtagtas
33301	cactgtgttg	gtgaaaaagc	acagctaaat	caaaagaaac	gcgattttca	aggegeeeaa
33361	cggtggcttc	caacaaagcc	tccacgcgca	catccaagaa	caaaagaata	ccaaaagaag
33421	gagcattttc	taactcctca	atcatcatat	tacattcctg	caccattccc	agataatttt
33481	cagctttcca	gccttgaatt	attcgtgtca	gttcttgtgg	taaatccaat	ccacacatta
33541	caaacaggtc	ccggagggcg	ccctccacca	ccattcttaa	acacaccctc	ataatgacaa
33601	aatatcttgc	tcctgtgtca	cctgtagcga	attgagaatg	gcaacatcaa	ttgacatgcc
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33901	aaaayaaaaa	titgecaaaa	adacacccaa ***********	attactotoggg	222224222	2222222
33961	cacacrácác	tccaacattg	ttagttttga	actagicige	aaaaataaaa	+
34021	gcgtcatatc	atagtagcct	gacgaacagg	tggataaatc	agtctttcca	tcacaayaca
34081	agccacaggg	tctccagctc	gaccctcgta	aaacctgtca	tggtgattaa	acaacagcac
34141	cgaaagttcc	tcgcggtgac	cagcatgaat	aattcttgat	gaagcataca	atccagacat
34201	gttagcatca	gttaacgaga	aaaaacagcc	aacatagcct	ttgggtataa	ttatgcttaa
34261	tcotaagtat	agcaaagcca	cccctcgcgg	atacaaagta	aaaggcacag	gagaataaaa
34321	aatataatta	tttctctgct	actattcagg	caacgtcgcc	cccggtccct	ctaaatacac
34381	atacaaagcc	tcatcagcca	taacttacca	gacaaagtac	agcgggcacg	cacaagctct
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345UT	aycadaycyt	aaaaaattttt	gccaaaccca	anduracet	+cc+c+++c+	cacggtacgt
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34621	cacateceat	taacttgcaa	cgtcattttc	ccaeggeege	geegeeegt	ttagccgtta
34681	accccacagc	caatcaccac	acaccccaca	atttttaaaa	tcacctcatt	tacatattgg
	caccattcca	tctataaggt	atattattga	rgatg		
	D NO: 12		,			

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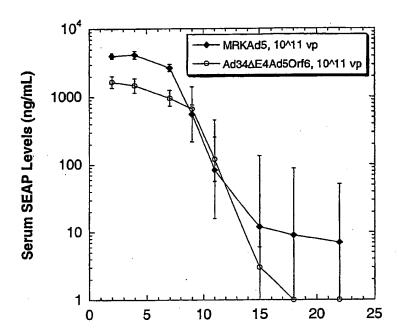


FIG. 29

Vaccine	Monkey	P	re	W	k 4	8	k B	W	24	W)	28	W	35
Wk 0, 4, 24	ID _	Mock	Gag*	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Gag
MRKAd5gag, 1041 vp MRKAd5gag, 1041 vp MRKAd5gag, 1041 vp	00C018 00C034 00C058	1 0 4	5 4 4	13 5 3	1025 219 1086	0 5 0	824 404 440	8 3 4	756 445 1439	0 3 0	474 339 2338	0 0 0	383 216 940
Ad34AE1gagAE4Ad5Orl6, 10^11 vp Ad34AE1gagAE4Ad5Orl6, 10^11 vp Ad34AE1gagAE4Ad5Orl6, 10^11 vp	00D038 00D042 00D066	6 6 3	8 30 18	5 4 1	111 89 118	1 4 1	301 264 816	0 1 0	224 73 429	1 0 0	535 181 439	0 0 0	233 69 273

Vaccine	Monk ID		D4 <sup>+</sup> CD3 <sup>+</sup> mphocytes	IFN-γ <sup>+</sup> CD8 <sup>+</sup> CD3 <sup>+</sup> per 10 <sup>6</sup> Lymphocytes			
	Ī	Mock	Gag <sup>a</sup>	Mock	Gag <sup>a</sup>		
Ad34∆E1gag∆E4Ad5Orf6	00D038	22	154	130	450		
	00D042	32	118	96	171		
	00D066	12	238	150	442		

Vancine	Veccine	Monkey	P	Ta .	T=4	wks	T≕Ð	wks	T=24	wke	T=26	wice	T=32	2 Wks
T=0, 4 w/cs	T=24 wks	ID	Mock	Gāy"	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Clag	Mock	Gag
Ad34AE1gagAE4Ad5Od6, 10*11 vp	Ad3S&E1gag&E4Ad5Orl8, 10^10 vp	00C016	4	8	1	84	5	334	5	29	0	308	3	244
Ad34AE1gagAE4Ad3Orf8, 10^11 vp	Ad35&E1grg&E4Ad5Od8, 10^10 vp	000044	1	1	8	79	0	374	В	138	٥	493	1 1	253
Ad34&E1gag&E4Ad5Od6, 10^11 vp	Ad356E1gag6E4Ad5Od6, 10^10 vp	00D064	4	6	1 1	125	В	655	6	145	٥	351	1	235
				L		Щ_							L	┷
Nzive		00D087	1	1	3	3	8	54	В .	B	5	5	3	0

Vaccine (T=0, 4 Wks) Vaccine (T=24 Wk)		Monkey		D4*CD3* mphocytes	IFN-7*CD8*CD3* per 10 <sup>8</sup> Lymphocytes		
		ID	Mock	Gag	Mock	Gag	
Ad34AE1gagAE4Ad5Orf6, 10^11 vp	Ad35AE1gagAE4Ad5Orf6, 10^10 vp	00D016	62	433	176	1288	
Ad34&E1gag&E4Ad5Orf6, 10^11 vp	Ad35AE1gagAE4Ad5Orf6, 10^10 vp	00D044	136	593	323	1871	
Ad34ΔE1gagΔE4Ad5Orf6, 10^11 vp	Ad35ΔE1gagΔE4Ad5On6, 10^10 vp	00D064	188	785	292	992	

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